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/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
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9-133-521-6
8-679-493A-158
9-230-371A-24
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7-792-988-2
8-649-006A-9
9-771-023-11
8-679-493A-159
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ALIGNMENTS

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RESULT 1
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US-08-147-492B-11
US-08-147-492B-11
APPLICANT: Hawkins, Phillip R.
APPLICANT: Bandaman, Oldse
CORRESPONDENGE ADDRESSED CHENOKINES, THEIR
TITLE OF INVENTION: EXPRESSED CHENOKINES,
TITLE OF INVENTION: EXPRESSED CHENOKINES,
TITLE OF INVENTION: PRODUCTION AND USES
ONDERSSEE: 12
CORRESPONDENGE ADDRESSE
APPLICANTS: 10-S
APPLICANTS: Pal-A ALC
STRET: 3174 PORTER DAI-B
COUNTRY: U.S.
ZIP: 94304
STRET: BM Compatible
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: Pal-NOV-1994
APPLICATION NUMBER: US/08/347,492B
FILING DATE: 02-NOV-1994
APPLICATION NUMBER: 08/103,011
FILING DATE: 05-OCT-1994
APPLICATION NUMBER: 33,954
APPLICATION NUMBER:
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US-08-347-492B-11

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APPLICANT: Wilde, Craig G.
APPLICANT: Wilde, Craig G.
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: ITS PRODUCTION AND USES
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
STREET: 3330 HILLVIEW AVENUE
CITY: PALO ALTO
STATE: CALIFORNIA
COUNTRY: USA
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Pred. No. 1.3e-50;
0; Mismatches 0;
Score 486; DB 1;
Pred. No. 1.3e-50;
); Mismatches 0;
                                                                                                                    ZIP: 94304

ZIP: 94304

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/375,346A
FILING DATE: 19-JAN-1995
CLASSIFICATION: 435
PRIOR APPLICATION 0ATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: LUTHER, BARBARA J.
REGISTRATION NUMBER: PF-0026 US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 855-0555
TELEPHONE: (415) 855-0572
                                                                                                                                                                                                                           Sequence 4, Application US/08375346A Patent No. 5605817 GENERAL INFORMATION:
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Similarity 100.0%;
92; Conservative C
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SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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ORIGINAL SOURCE:
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SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/421,144A
FILING DATE: 13-APR-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33954
REFERENCE/DOCKET NUMBER: PF-0031 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-855-0195
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1.3e-50;
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Pred. No. 1.3(); Mismatches
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APPLICANT: Hawkins, Phillip R.
APPLICANT: Bandman, Olga
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: EXPRESSED CHEMOKINES,
TITLE OF INVENTION: PRODUCTION AND USES
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                                                                   APPLICANT: BANDMAN, OLGA
APPLICANT: COLEMAN, ROGER
APPLICANT: STUART, SUSAN G.
TITLE OF INVENTION: NEW CHEMOKINE EXPRESS
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
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Patent No. 5936068
                                                                                                   US-08-421-144A-4
; Sequence 4, Application US/08421144A
; Patent No. 5874211
; GENERAL INFORMATION:
; APPLICANT: BANDMAN, OLGA
; APPLICANT: COLEMAN, ROGER
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Palo Alto
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TYPE: amino acid
STRANDEDNESS: single
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GENERAL INFORMATION:
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Pred. No. 1.3e-50;
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-08-467-123B-4
Sequence 4, Application US/08467123B
Sequence 4, Application US/08467123B
Sequence 4, Application US/08467123B
Sequence 4, Application:
Setent No. 5945506
GENERAL INFORMATION:
APPLICANT: Coleman, Roger
APPLICANT: Wilde, Craig C.
APPLICANT: Wilde, Craig C.
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: CHEMOKINE EXPRESSED IN
TITLE OF INVENTION: ITS PRODUCTION AND USE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 PAVVFQTKRSKQVCADPSESWVQEYVYDLELN
61 PAVVFQTKRSKQVCADPSESWVQEYVYDLELN
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
CITY: Palo Alto
STATE: CA
                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: Fast SEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/798,143
FILING DATE: 10-FEB-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: 08/347,492
FILING DATE: 29-NOV-1994
APPLICATION NUMBER: 08/320,011
FILING DATE: 05-OCT-1994
APPLICATION NUMBER: 08/320,011
FILING DATE: 05-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0024
TELEPHONE: 415-855-0555
TELEPHONE: 415-852-0195
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 2014
ADDRESSEE: Incyte Pharmaceuticals,
STREET: 3174 Porter Drive
CITY: Palo Alto
STATE: CA
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I 127080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: sin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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                                                                                       94304
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Best Local S
Watches 92
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1 MKLCVTVLSLLMLVAAFCSPALSAPMGSDPPTACCFSYTARKLPRNFVVDYYETSSLCSQ
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APPLICANT: Swanberg, Stephen
TITLE OF INVENTION: CHIMERIC POLYPEPTIDES CONTAINING
TITLE OF INVENTION: CHEMOKINE DOMAINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/808,720
FILING DATE:
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 486; DB 2;
Pred. No. 1.3e-50;
; Mismatches 0;
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                                                                                                                                                                                                                                                                       PF-0026-1 DIV
                                                                                                                                                                        APPLICATION NUMBER: US 08/375,346
FILING DATE: 19-JAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0026-1 I
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-555-0555
TELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Genetics Insititute, Inc
STREET: 87 CambridgePark
CITY: Cambridge
                                                                                                        US/08/467,123B
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,1
FILING DATE: 06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P-41,323
ER: GI5291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 10, Application US/08808720
Patent No. 6100387
GENERAL INFORMATION:
APPLICANT: Herrmann, Steve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Sprunger, Suzanne
REGISTRATION NUMBER: P-41,323
REFERENCE/DOCKET NUMBER: G152
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8284
TELEFAX: (617) 876-5851
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                                                                                                                                        CLASSIFICATION: 530
PRIOR APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: sing
TOPOLOGY: linear
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INFORMATION FOR
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STATE:
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Best Local S
Matches 92
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Best Local &
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ORGANISM:
                                                                                                                                                  Best Local
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US-08-6
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Patent No. 6281200
GENERAL INFORMATION:
APPLICANT: Fife, Kenneth H.
APPLICANT: Krathwohl, Mitchell D.
APPLICANT: Hromas, Robert
APPLICANT: Brown, Darron R.
APPLICANT: Brown, Darron R.
APPLICANT: Brown CHNCTIONAL CHARACTERIZATION OF THE C-C CHEMOKINE-LIKE
TITLE OF INVENTION: MOLECULES ENODED BY MOLLUSCUM CONTAGIOSUM VIRUS TYPES 1
TITLE OF INVENTION: AND 2
TITLE OF INVENTION: AND 2
FILE REFERENCE: INDY:034
CURRENT APPLICATION NUMBER: 60/055,532
EARLIER FILING DATE: 1999-08-18
EARLIER FILING DATE: 1997-08-15
NUMBER OF SEQ ID NOS: 30
SOFTWARE: PatentIN Ver. 2.0
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1.3e-50;
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Pred. No. 1.3e-50;
0; Mismatches 0;
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3-08-679-493A-157
Sequence 157, Application US/08679493A
Patent No. 6303295
GENERAL INFORMATION:
APPLICANT: Taylor, Ethan W.
TITLE OF INVENTION: SELENOPROTEINS, CODING SEFILE REFERENCE: 55-95
CURRENT APPLICATION NUMBER: US/08/679,493A
CURRENT FILING DATE: 1996-07-12
PRIOR APPLICATION NUMBER: 60/001203
PRIOR FILING DATE: 1995-07-14
                                                                                                                                   ed. No. 1.
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100.0%; Pred. No. 3
tive 0; Mismatche
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SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
TYPE: amino acid
STRANDEDNESS:
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92; Conservative
                                                                                                                                               Conservative
                                                                     linear
: protein
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                                                                                                                                   Similarity
                                                                   TOPOLOGY: li
MOLECULE TYPE:
3-08-808-720-10
                                                                                                                                             92;
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        INFORMATION
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APPLICANT: Chang, Yuan
APPLICANT: Bohenzky, Roy A
APPLICANT: Moore, Patrick S
APPLICANT: Moore, Patrick S
TITLE OF INVENTION: UNIQUE ASSOCIATED KAPOSI'S SARCOMA VIRUS SEQUENCES AND TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: 45185-G-PCT-US
CURRENT APPLICATION NUMBER: US/09/230,371A
CURRENT FILING DATE: 1999-11-17
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Sequence 158, Application US/08679493A
Patent No. 6303295
GENERAL INFORMATION:
APPLICANT: Taylor, Ethan W.
TITLE OF INVENTION: SELENOPROTEINS, CODING SEQUENCES AND METHODS
FILE REFERENCE: 55-95
                                                                                                                                                                                        Length 92;
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1.3e-50;
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Pred. No. 1.3e-50;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                        PAVVEQTKRSKQVCADPSESWVQEYVYDLELN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/08/679,493A CURRENT FILING DATE: 1996-07-12 PRIOR APPLICATION NUMBER: 60/001203 PRIOR FILING DATE: 1995-07-14 PRIOR APPLICATION NUMBER: 60/003,112 PRIOR FILING DATE: 1995-09-01 NUMBER OF SEQ ID NOS: 216 SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24, Application US/09230371A
Patent No. 6348586
GENERAL INFORMATION:
PRIOR APPLICATION NUMBER: 60/003,11:
PRIOR FILING DATE: 1995-09-01
NUMBER OF SEQ ID NOS: 216
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 157
LENGTH: 92
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US-08-679-493A-157
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92; Conserv
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Swanberg, Stephen
VENTION: CHIMERIC POLYPEPTIDES CONTAINING
VENTION: CHEMOKINE DOMAINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/808,720
FILING DATE:
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Sprunger, Suzanne
REGISTRATION NUMBER: P-41,323
REFERENCE/DOCKET NUMBER: G15291
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (617) 498-8284
TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 486; DB 3;
Pred. No. 6.4e-50;
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Pred. No. 1.3e-50;
Mismatches 0
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3-08-808-720-7
Sequence 7, Application US/08808720
Patent No. 6100387
GENERAL INFORMATION:
APPLICANT: Herrmann, Steve
APPLICANT: Swanberg, Stephen
TITLE OF INVENTION: CHIMERIC POLYPEPTIDE
TITLE OF INVENTION: CHEMOKINE DOMAINS
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Insititute, Inc.
STREET: 87 CambridgePark
CITY: Cambridge
STATE: MA
    PCT/US97/13346
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PRIOR APPLICATION NUMBER: PCT
PRIOR FILING DATE: 1997-07-22
NUMBER OF SEQ ID NOS: 30
SOFTWARE: Patentin Ver. 2.0
EQ ID NO 24
LENGTH: 92
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100.0%;
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amino acid
                                                                                                                                                               Similarity 100
12; Conservative
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NECULE TYPE: protein
108-720-7
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                                                                                                      ORGANISM: Human
-09-230-371A-24
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Best Local S
Matches 92
                                                                                        TYPE: PRT
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Best Local (
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1 MKLCVTVLSLLMLVAAFCSPALSAPMGSDPPTACCFSYTARKLPRNFVVDYYETSSLCSQ 60
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Sequence 159, Application US/08679493A

Patent No. 6303295

GENERAL INFORMATION:
APPLICANT: Taylor, Ethan W.

TITLE OF INVENTION: SELENOPROTEINS, CODING SEQUENCES AND METHODS
FILE REFERENCE: 55-95

CURRENT APPLICATION NUMBER: US/08/679,493A

CURRENT FILING DATE: 1996-07-12

PRIOR APPLICATION NUMBER: 60/001203

PRIOR FILING DATE: 1995-09-01
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Pred. No. 3.9e-50;
0; Mismatches 1;
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ed. No. 3.9e-50;
Mismatches 1
               61 PAVVFQTKRSKQVCADPSESWVQEYVYDLELN 92
PAVVFQTKRSKQVCADPSESWVQEYVYDLELN 92
                                                                                                                                                  APPLICANT: Hayward, Gary
APPLICANT: Hayward, Gary
APPLICANT: Nicholas, John
APPLICANT: Hardwick, J. Marie
APPLICANT: Hardwick, J. Marie
APPLICANT: Reitz, Marvin
TITLE OF INVENTION: No. 6264958el Genes of Ka
TITLE OF INVENTION: Associated Herpesvirus
FILE REFERENCE: 1107.78372
CURRENT APPLICATION NUMBER: US/09/230,637
CURRENT FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: 60/022,591
PRIOR FILING DATE: 1996-07-25
PRIOR FILING DATE: 1997-07-24
NUMBER OF SEQ ID NOS: 62
SOFTWARE: FastSEQ for Windows Version 4.0
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Pred. No. 3
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SOFTWARE: Patentin Ver. 2.0
LENGTON:
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Patent No. 6264958
GENERAL INFORMATION:
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larity 98.9%;
Conservative (
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US-09-230-637-41
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                                                                                             US-09-230-637-41
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SOFTWARE: Pate
SEQ ID NO 159
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macrophage inflammatory
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Pred. No. 2e-49;
0; Mismatches
                                                                 ESULT 14
3-07-792-988-2
Sequence 2, Application US/07792988
Patent No. 5306709
GENERAL INFORMATION:
APPLICANT: Gewirtz, Alan M.
TITLE OF INVENTION: Suppression of megakary
TITLE OF INVENTION: Cytopoiesis by macrophe
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: University of Pennsylvania
STREET: Suite 419
STREET: Suite 419
STREET: Suite 419
STREET: U.S.A.
ZIP: Philadelphia
STATE: Pennsylvania
COUNTRY: U.S.A.
ZIP: 19104-3246
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 720 Kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         92
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MEDIUM TIFE.

COMPUTER: IBM PS/2
OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/792,988
FILING DATE: 19911115
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Monaco, Daniel A.
REGISTRATION NUMBER: 6056-159
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-8383
TELEFAX: (215) 568-8383
TELEFAX: (215) 568-5549
TELEX: NO. 5306709e
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
TWORM AMINO ACID
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-08-649-006A-9

Sequence 9, Application US/08649006A

Patent No. 6548654

GENERAL INFORMATION:

APPLICANT: Hardiman, Gerard T.

APPLICANT: Rossi, Devora L.

APPLICANT: Bacon, Kevin B.

APPLICANT: Bacon, Cernando
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Similarity 98.9%;
91; Conservative
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TYPE: AMINO ACID
STRANDEDNESS: single st
TOPOLOGY: linear
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Best Local
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Gaps
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/649,006A

FILING DATE: 16-MAY-1996

CLASSIFICATION DATA:

APPLICATION NUMBER: US 08/590,828

FILING DATE: 24-JAN-1996

ATTORNEY/AGENT INFORMATION:

NAME: Ching, Edwin P.

REGISTRATION NUMBER: 34,090

REFERENCE/DOCKET NUMBER: DX0569K

TELECOMMUNICATION INFORMATION:

TELEPHONE: (650)852-9196

INFORMATION FOR SEQ ID NO: 9:
APPLICANT: Schall, Thomas J.
APPLICANT: Zlotnik, Albert
TITLE OF INVENTION: Mammalian CX3C Chemokine Genes
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: DNAX Research Institute
STREET: 901 California Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94.9%; Score 461; DB 4; I larity 92.4%; Pred. No. 1.2e-47; Conservative 4; Mismatches 3;
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LENGTH: 92 amino acids
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California
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STRANDEDNESS: not
TOPOLOGY: linear
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85; Conserv
                                                                                                                                                          COUNTRY: USA
ZIP: 94304-1104
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Matches 85
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5.1.6 Compugen Ltd GenCore version (c) 1993 - 2003 Copyright

using sw model - protein search, protein

2003, 15:36:20 ; Search time 13.4262 Seconds (without alignments) 651.810 Million cell updates/se December 16, : uo d

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91

US-09-920-137A-8 478 :le:
rfect score:

......VCANPEKKWVREYINSLEMS 1 MKVSAARLAVILIATALCAP. inence:

oring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

283308 segs, 96168682 residues rched:

283308

hits satisfying chosen parameters: of al number

iimum DB seq length: 0
iimum DB seq length: 2000000000

t-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR 76:*
1: pir1:*
2: pir2:*
3: pir3:* ч си се 4

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

STIMMADIES

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SUMMARIES		2881	4653	4673	3567	3057	5232	3055	3239	3176	0491	5467	4814	0772	2529	2213	5040	1809	A30209	3017	3929	2233	1241	1809	2247	5029	1685	955	HUL	242
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ALIGNMENTS

RESULT 1	
A28815	
Monocyte chemoattractant cytokine RANTES precursor - human N;Alternate names: small inducible cytokine A5: T-cell specific cytokine pantes	
C;Date: 30-Jun-1989 #sequence_revision 16-Aug-1996 #text_change 29-May-1998	
J. Immunol. 141, 1018-1025, 1988	4
A: Title: A human T cell-specific molecule is a member of a new gene family.	
A; Note: Felice number: Azasıs; Moid: 88285659; PMID: 2456327 A; Accession: A28815	
A; Molecule type: mRNA	
A;Residues: 1-91 <sch></sch>	
pon Activation,	E NO N
A; Gene: GDB:SCYA5; D17S136E	
A;Cross-references: GDB:120749; OMIM:187011	
A; Map position: 17q11.2-17q12	
C; Superfamily: macrophage inflammatory protein	
Vineywords: cnemotaxis; cytokine; immediate-early protein; inflammation; T-cell F:1-23/Domain: gional semience #status prodicted off:	
F;24-91/Product: T-cell protein RANTES #status predicted <mat></mat>	
Query Match 98.7%; Score 472; DB 1: Length 91:	
ocal Similarity 98.9%; Pred. No. 2.4e-44; s 90; Conservative 0; Mismatches 1: Indels	
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I MANSAAKLAVILLAID	
DD 1 MKVSAAALAVILIATALCAPASASPYSSDTTPCCFAYIARPLPRAHIKEYFYTSGKCSNP 60	
QY 61 AVVFVTRKNRQVCANPEKKWVREYINSLEMS 91	
DD 61 AVVFVTRKNRQVCANPEKKWVREYINSLEMS 91	

#text_change 22-Jun-1999 monocyte chemoattractant cytokine RANTES precursor - mouse monocyte chemoattractant cytokine RANTES precursor - mouse N; Alternate names: MuRantes C; Species: Mus musculus (house mouse) C; Date: 18-Jun-1993 #sequence revision 16-Aug-1996 #text_change 2; C; Date: 18-Jun-1993 #sequence revision 16-Aug-1996 #text_change 2; C; Accession: I48875; A46539; I48654; I56970
B; Danoff, T.M.; Lalley, P.A.; Chang, Y.S.; Heeger, P.S.; Neilson, J. Immunol. 152, 1182-1189, 1994
A; Title: Cloning, genomic organization, and chromosomal localization; A; Reference number: I48875; MUID: 94132613; PMID: 7507961
A; Accession: I48875
A; Accession: I48875
A; Accession: Lype: DNA

localization of

gene

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the

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macrophage inflammatory protein 1-alpha precursor - human
N;Alternate names: LD78-alpha protein precursor; lymphocyte tumor promoter-induced prot
tivation protein 1
C;Species: Homo sapiens (man)
C;Date: 03-Aug-1992 #---
1 MKVSAARLAVILIATALCAPASASPYSSDT-TPCCFAYIARPLPRAHIKEYFYTSGKCSN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 MOVSTAALAVLLCTMALCNOVLSAPLAADTPTACCFSYTSROIPONFIADYFETSSOCSK
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:: LD78-beta protein #status predicted
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Mismatches
                                                  60 PAVVFVTRKNRQVCANPEKKWVREYINSLEMS
                                                                    61 PAVVFQTKKGRQVCANPSESWVQEYVDDLELN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Introns: 26/1; 64/2
C;Superfamily: macrophage inflammatory protein
C;Keywords: cytokine
F;1-22/Domain: signal sequence #status predict
F;23-93/Product: LD78-beta protein #status pred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Cross-references: GDB:120369; OMIM:182284
A; Map position: 17q11-17q21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25;
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                  PID:9460091
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                                                                                                                                                                                NCBIP:106770)
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                                                                                       dolecule type: mRNA
Residues: 1-18,'A',20-91 <SCH>
Cross-references: GB:S37648; NID:g250207; PIDN:AAB22330.1; PID:g250208
Experimental source: macrophage cell line PU5-1.8
Vote: sequence extracted from NCBI backbone (NCBIN:106768, NCBIP:106776)
Shin, H.S.; Drysdale, B.E.; Shin. M.T. MACLIT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           1-40,'E',42-91 <NEI>
1-40,'E',42-91 <NEI>
2rences: GB:M77747; NID:g200649; PIDN:AAA40029.1; PID:g200650
this chemoattractant for monocytes but not neutrophils is an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Σ
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                                                                cytokine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Introns: 26/1; 63/2
Superfamily: macrophage inflammatory protein
Seywords: chemotaxis; cytokine; immediate-early protein; inflammation
--23/Domain: signal sequence #status predicted <SIG>
-4-91/Product: monocyte chemoattractant cytokine RANTES #status predic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      S.; Yoshinaga,
                                                                                                                                                                                                                                 in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lori, S.; Goto, K.; Goto, F.; Mutakami, K.; Ohkawara, S.; Yoshinaga,
   Immunol. 6, 149-156, 1994
litle: Dynamic changes in mRNA expression of neutrophils during the
eference number: 146730; MUID:94198229; PMID:8148323
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                                                                 RANTES
                                                                                                                                                                                                                                 element
               PIDN: AAA18302.1;
                                                                                                                                                                                                                                                                           Status: translation not shown; translated from GB/EMBL/DDBJ
folecule type: DNA
Residues: 1-91 <SHI>
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                                                              of the murine
PMID:1376260
                                                                                                                                                                                         n, H.S., Drysdale, B.E., Shin, M.L., Noble, P.W., Cell. Biol. 14, 2914-2925, 1994
le: Definition of a lipopolysaccharide-responsive erence number: 148654; MUID: 94217689; PMID: 7513046
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nes 28;
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Pred. No. 9.7e-36;
7; Mismatches 12;
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               460090;
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Tross-references: EMBL:U02298; NID:g46009
Schall, T.J.; Simpson, N.J.; Mak, J.Y.
r. J. Immunol, 22, 1477-1481, 1992
Fitle: Molecular cloning and expression caference number: A46539; MUID:92289805;
                                                                                                                                                                                                                                                                                                                                            Wellson, E.G.; Krensky, A. Int. 41, 220-225, 1992
Title: Isolation and characterization of Seference number: I56970; MUID:92277990; Accession: I56970
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ss: Oryctolagus cuniculus (domes
14-Feb-1997 #sequence_revision
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llarity 79.1%;
Conservative
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litle: Definition
leference number:
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lomment: This chemienetics:
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Cross-references: GB:D90144; NID:g219905; PIDN:BAA14172.1; PID:g219906
Zipfel, P.F.; Balke, J.; Irving, S.G.; Kelly, K.; Siebenlist, U.
Immunol. 142, 1582-1590, 1989
Title: Mitogenic activation of human T cells induces two closely related
Reference number: A30574; MUID:89140347; PMID:2521882
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Title: Three human homologs of a murine gene encoding an inhibitor of Reference number: A30412; MUID:91103879; PMID:2271120
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Accession: A30412
Molecule type: mRNA
Residues: 1-92 <BLU>
Cross-references: GB:M23178; GB:M32337; NID:g182846; PIDN:AAA35858.1;
Cross-references: GB:M23178; GB:M32337; NID:g182846; PIDN:AAA35858.1;
Obaru, K.; Fukuda, M.; Maeda, S.; Shimada, K.
Biochem. 99, 885-894, 1986
Title: A cDNA clone used to study mRNA inducible in human tonsillar ly
Reference number: A24198; MUID:86223879; PMID:3086300
Accession: A24198
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Accession: A35673; A30574; A30412; A24198; A30908
Nakao, M.; Nomiyama, H.; Shimada, K.
1. Cell. Biol. 10, 3646-3658, 1990
Title: Structures of human genes coding for cytokine l
Reference number: A35673; MUID:90287155; PMID:1694014
Accession: A35673
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Wolecule type: mRNA
Residues: 1-92 <RES>
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Shi, M.M.; Godleski, J.J.; Paulauskis, J.D.
Schem. Biophys. Res. Commun. 211, 289-295, 1995
Fitle: Molecular cloning and posttranscriptional
Reference number: I52322; MUID:95298037; PMID:777
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Map position: 17q11-17q21
Superfamily: macrophage inflammatory protein
1-20/Domain: signal sequence #status predicted
21-92/Product: macrophage inflammatory protein
33-57,34-73/Disulfide bonds: #status predicted
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Species: Rattus norvegicus (Norway ra
Date: 29-May-1998 #sequence_revision
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Cross-references: GB: 3enetics: 3ene: GDB:SCYA3
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Cross-references:
Blum, S.; Forsdyke
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Molecule type: mR
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Molecule type:
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Macrophage inflammatory protein 1-beta precursor - mouse
Macrophage inflammatory protein 1-beta precursor - mouse
Natherate chase: 1440; S125 gamma; T-cell activation protein gamma
C_Species Mus musculus (buse mouse)
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T_mmunol. 142, 679-687, 1969
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        Score 248; DB 2;
Pred. No. 5.9e-20;
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ilarity 45.7%;
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51.9%;
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SIS /MIP-1a; crophage inflammatory protein-1-alpha precursor - mouse Alternate names: heparin-binding chemotaxis protein; L2G25B protein; SCI Species: Mus musculus (house mouse)
Date: 17-Jul-1992 #sequence revision 17-Jul-1992 #text change 16-Jul-1999 Accession: S11685; A32393; S04533; A53885; A30552; PS0303; A27596; I5610 Grove, M.; Lowe, S.; Graham, G.; Pragnell, I.; Plumb, M. cleic Acids Res. 18, 5561, 1990
Title: Sequence of the murine haemopoietic stem cell inhibitor/macrophag Reference number: S11685; MUID: 91016858; PMID: 2216738

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e inflammatory

Accession: \$11685 Molecule type: DN

DNA

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ammatory prot Residues: 1.92 <GRO>
Cross-references: EMBL:X53372; NID:g54062; PIDN:CAA37452.1; PID:g297531
Note: the authors' translation of the nucleotide sequence differs at several
Note: the authors' translation of the nucleotide sequence differs at several
Kwon, B.S.; Weissman, S.M.
OC. Natl. Acad. Sci. U.S.A. 86, 1963-1967, 1989
Title: cDNA sequence of two inducible T-cell genes.
Reference number: A32393; MUID:89184547; PMID:2784565
Accession: A32393
Molecule type: mRNA
Residues: 1-92 <KWO>
Cross-references: GB:J04491; NID:g201524; PIDN:AAA40304.1; PID:g201525
Davatelis, G.; Tekamp-Olson, P.; Wolpe, S.D.; Hermsen, K.; Luedke, C.; Galle
Exp. Med. 167, 1939-1944, 1988
Title: Cloning and characterization of a cDNA for murine macrophage inflamma

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S.D.; Wolpe, Title: Ctoning and Characterization of a CDNA for mutine macrophage inflamma Accession: 80453; MUID:88258380; PMID:3290382
Accession: 804533
Molecule type: mRNA
Residues: 1-46, E'. 50-90, I'. 92 cDA2>
Cross-references: EMBL:X12531
Note: the authors translated the codon GAG for residue 49 as Asp and ATT for Note: the sequence has been corrected in reference A53885
Davatelis, G.; Tekamp-Olson, P.; Wolpe, S.D.; Hermsen, K.; Luedke, C.; Galle Exp. Med. 170, 2189, 1989
Note: the sequence has been corrected in reference A53885
Davatelis, G.; Tekamp-Olson, P.; Wolpe, S.D.; Hermsen, K.; Luedke, C.; Galle Exp. Med. 170, 2189, 1989
Contents: erratum
Accession: A53885
Contents: erratum
Accession: A53885
Contents: EMBL:X12531; NID:953122; PIDN:CAA31047.1; PID:953123
Reference number: A53885
Cross-references: EMBL:X12531; NID:953122; PIDN:CAA31047.1; PID:953123
Rown, K.D.; Zurawski, S.M.; Mosmann, T.R.; Zurawski, G.
Immunol. 142, 679-687, 1989
Accession: A30552; MUID:89093958; PMID:2521353
Accession: A30552; MUID:89093958; PMID:2521353
Accession: A30552; MUID:89093958; PMID:2521353
Accession: A30552; MUID:89093958; PMID:3AA40146.1; PID:9533241
Sherry, B.; Tekamp-Olson, P.; Gallegos, C.; Bauer, D.; Davatelis, G.; Wolpe, Exp. Med. 168, 225-2259, 1988
Accession: PS0303
Accession: PS0303
Accession: PS0303
Accession: PS0303
Accession: PS0303; PMID:89067830; PMID:3058856
Accession: PS0303; ACCESSION AC

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le type: mRNA es: 24-33,'XX',36-54 <SHE> S.D.; Davatelis, G.; Sherry, B.; Beutler, B.; Hesse, D.G.; Nguyen, H.T.; M Med. 167, 570-581, 1988 Macrophages secrete a novel heparin-binding protein with inflammatory and Macrophages secrete a novel heparin-binding protein with inflammatory and , S.D.; Dav Med. 167,

.; Э Sherry, were also found er, S.; Manogue, K.R.;

and conserv of murine macrophage inflammatory protein-1-alpha MUID:91237116; PMID:2033269 Accession: I56104
Status: preliminary; translated from GB/EMBL/DDBJ
Molecule type: DNA
Residues: 1-92 <RES> Title: Macrophysical Reference number: A27596; MUID: BOLDANDACCESSION: A27596
Molecule type: protein Residues: 24-33,'XX',36-42 <WOLDANDE: 26-Met, 30-Pro, and 39-Thr were Widmer, U.; Yang, Z.; van Deventer, S. Immunol. 146, 4031-4040, 1991
Title: Genomic structure of murine macalline residues: I56104; MUID: 9123711

GB:M73061; NID:g199694; PIDN:AAA39707.1; PID:g199695 Cross-references:

C;Comment: This protein is a monokine.
C;Genetics:
A;Introns: 23/3; 26/1; 63/2
A;Introns: 23/3; 26/1; 63/2
C;Superfamily: macrophage inflammatory protein
C;Keywords: heparin binding
C;Keywords: heparin binding
F;1-23/Domain: signal sequence #status predicted <SIG>
F;24-92/Product: macrophage inflammatory protein #status experimental

Gaps Length 92; Indels 26; Score 243; DB 2; Pred. No. 2.1e-19; Mismatches 19; 50.8%; Conservative Similarity 45; Query Match Best Local

2;

59 1 MKVSAARLAVILIATALCAPASASPYSSDT-TPCCFAYIARPLPRAHIKEYFYTSGKCSN

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Macrophage inflammatory protein 1-beta precursor [validated] - human
Marcophage inflammatory protein 1-beta precursor [validated] - human
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Charcesion, Mainty Add 78, 7-cell activation protein gamma
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A Accession (Mainty Add 78, 7-cell activation gene (LAG-1).

A Residues: 1-92 - CAN
A Residues: 1-94 - CAN
A Residues: 1-14 - CAN
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             MKVSAALLWILLIAAAFSPQGLTGPASVPTT-CCFNLANRKIPLQRLESYRRITSGKCPQ
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C;Species: Cavia porcellus (guinea pig)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text
C;Accession: 148147
R;Yoshimura, T.
J. Immunol. 150, 5025-5032, 1993
A;Title: cDNA cloning of guinea pig monocyte chemoattrac
A;Reference number: 148147; MUID:93267104; PMID:8496603
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 167; DB 2; L
Pred. No. 4.5e-11;
3; Mismatches 34;
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71 REAVIFKTKLDKEICADPTQKWVQDFMKHLD 101
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                                                                          60 PAVVFVTRKNROVCANPEKKWVREYINSLE
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Date: 01-Nov-1996 #sequence_revision 01-Nov-1996 #text_change 20-Jun-2000
Accession: JC4912
Bartels, J.; Schlueter, C.; Richter, E.; Noso, N.; Kulke, R.; Christophers, ochem. Biophys. Res. Commun. 225, 1045-1051, 1996
Title: Human dermal fibroblasts express eotaxin: Molecular cloning, mRNA expecsion: JC4912
Reference number: JC4912; MJD:96374440; PMID:8780731
Accession: JC4912
Reference number: JC4912; MJD:96374440; PMID:8780731
Accession: JC4912
Reference number: JC4912; MJD:96374440; PMID:8780731
Crossion: JC4912
Residues: J-97 < BAR>
Cross-references: EMBL:Z75668; NID:91531982; PIDN:CAA99997.1; PID:91531983
Experimental source: dermal fibroblast
Comment: This protein has eosinophil specific chemotatic activity.
Superfamily: macrophage inflammatory protein
Keywords: fibroblast
1-18/Domain: signal sequence #status predicted <SIG>
1-18/Domain: signal sequence #status predicted <MAT>
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Map position: 17q21-17q21

Introns: 26/1; 64/2

Superfamily: macrophage inflammatory protein

Keywords: chemotaxis; cytokine; inflammation

1-23/Domain: signal sequence #status predicted <SIG>
24-92/Product: macrophage inflammatory protein 1-beta #status experimental
34-58;35-74/Disulfide bonds: #status experimental
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Accession: D30552
Molecule type: mRNA
Residues: 1-39, 'REASS', 46-92 <BRO>
Cross-references: GB:M23502; NID:g533212; PIDN:AAA36656.1; PID:g533213
Clore, G.M.; Lodi, P.J.; Garrett, D.S.; Gronenborn, A.M.
Clore, G.M.; Lodi, P.J.; Garrett, D.S.; Gronenborn, A.M.
Dmitted to the Brookhaven Protein Data Bank, January 1994
Reference number: A52206; PDB:1HUM
Reference number: A52206; PDB:1HUM
Contents: annotation; conformation and disulfide bond assignments by (1)
Comment: This protein is secreted by activated lymphocytes and monocytes
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                                                                                          Molecule type: mRNA
Residues: 7-55,'I',57-79,'T',81-92 <MIL>
Residues: 7-55,'I',57-79,'T',81-92 <MIL>
Cross-references: GB:M57503; NID:g339726; PIDN:AAA36752.
Brown, K.D.; Zurawski, S.M.; Mosmann, T.R.; Zurawski, G.
Immunol. 142, 679-687, 1989
Title: A family of small inducible proteins secreted by of various activation processes.
Reference number: A30552; MUID:89093958; PMID:2521353
Accession: D30552
                                      activated human prints PMID:2809212
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Pred. No. 3.2e-11;
16; Mismatches 35;
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Pred. No. 2.3e-19;
3; Mismatches 26;
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PAVVFQTKRSKQVCADPSESWVQEYVYDLELN
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Lumunol. 143, 2907-2916, 1989
Title: A novel polypeptide secreted by ac Reference number: A45817; MUID:90038522; Accession: B45817
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ilarity 41.1%;
Conservative 16
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C.Species: Sus scrofa domestica (domestic pig)
C.Accession: Sep-1993 #sequence_revision 20-Aug-1994 #text_change 16-Jul-1999
C.Accession: JC2136; S57498
R.Hosang, K.; Knoke, I.; Klaudiny, J.; Wempe, F.; Wuttke, W.; Scheit, K.H.
Biochem. Blophys. Res. Commun. 1999; 962-968, 1994
A.;Title: Porcine luteal express monocyte chemattractant protein-1 (MCP-1): Ana A; Reference number: JC2136; MUD:94183284; PMID:7510962
A.;Accession: JC2136; MUD:94183284; PMID:7510962
A.;Accession: JC2136; MUD:94183284; PMID:7510962
A.;Accession: S57497
A.;Residues: 1-99 +4MOS
A.;Cross references: GB:Z48479; NID:g837116; PIDN:CAA88370.1; PID:g872313
A.;Cross reference: manay
A.;Cross reference: EMBL:X79416; NID:g872312; PIDN:CAA55945.1; PID:g872313
C;Superfamily: macrophage inflammatory protein
C;Keywords: glycoprotein
F;1-23/Domain: signal sequence #status predicted <AMT>F;24-99/Product: monocyte chemoattractant protein-1 #status predicted
F;24-99/Product: monocyte chemoattractant protein-1 #status predicted
F;34/Binding site: carbohydrate (Asn) (covalent) #status predicted
                                                                                                  A; Molecule type: mRNA
A; Residues: 1-99 < VAN>
A; Residues: 1-99 < VAN>
A; Cross-references: GB: Y10802; NID: g1924937; PIDN: CAA71760.1; PID: g1924938
A; Cross-references: bone marrow
C; Comment: This protein belongs to the beta-chemokine family which is one of the major tis and in tumor biology, and contribute to the trafficking and recuitment of the resp
                           expression
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C;Superfamily: macrophage inflammatory protein
F;1-23/Domain: signal sequence #status predicted <SIG>
F;24-99/Product: monocyte chemotactic protein-2 #status predicted
Biochem. Biophys. Res. Commun. 231, 726-730, 1997
A;Title: Human monocyte chemotactic protein-2: cDNA cloning and
A;Reference number: JC5295; MUID:97224420; PMID:9070881
A;Accession: JC5295
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Pred. No. 8.4e-10;
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1.8e-09;
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Pred. No.
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37.4%; Pre
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34.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nmediate-early serum-responsive protein JE precursor - rat; Alternate names: monocyte chemoattractant protein-1; Species: Rattus norvegicus (Norway rat); Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-19; Accession: $07723; JN0128; Timmers, H.T.M.; Pronk, G.J.; Bos, J.L.; van der Eb, A.J. acleic Acids Res. 18, 23-34, 1990; Title: Analysis of the rat JE gene promoter identifies an AP-1 binding; Reference number: $07723; MUID: 90174947; PMID:2106664
                                                                                                         PID:9349821
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Cross-references: GB:M57441; NID:g205333; PIDN:AAA63496.1; PID:g20533.; Experimental source: spleen cells; Note: the authors translated the codon GAA for residue 62 as Lys and code.
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Superfamily: macrophage inflammatory protein
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; Residues: 1-148 <TIM>
; Residues: 1-148 <TIM>
; Cross-references: EMBL:X17053; NID:g55530; PIDN:CAA34901.1; PID:g55; Yoshimura, T.; Takeya, M.; Takahashi, K.; Yoshimura, T.; Takeya, M.; Takahashi, K.; iochem. Biophys. Res. Commun. 174, 504-509, 1991
; Title: Molecular cloning of rat monocyte chemoattractant protein-1; Title: Molecular cloning of rat monocyte chemoattractant protein-1; Reference number: JN0128; MUID:91128376; PMID:1704226
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larity 34.0%; Pred. No. 9.6e-10;
Conservative 22; Mismatches 36
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1; Mismatches 33
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SUMMARIES

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Hoover D.M., Shaw J., Gryczynski Z., Proudfoot A.E.I., Wells T.N.C., Lubkowski J.;

Lubkowski J.;

"The crystal structure of Met-RANTES: comparison with native RANTES and AOP-RANTES.";

Protein Pept. Lett., 7:73-82(2000).

Protein Pept. Lett., 7:73-82(2000).

-!- FUNCTION: CHEMOATTRACTANT FOR BLOOD MONOCYTES, MEMORY T HELPER CELLS AND EOSINOPHILS. CAUSES THE RELEASE OF HISTAMINE FROM BASOPHILS AND ACTIVATES EOSINOPHILS. BINDS TO CCR1, CCR3, CCR4 CCR5. ONE OF THE MAJOR HIV-SUPPRESSIVE FACTORS PRODUCED BY CD8+CCR5. ONE OF THE MAJOR HIV-SUPPRESSIVE FACTORS PRODUCED BY CD8+CCR5. ONE OF THE MAJOR HIV-SUPPRESSIVE FACTORS PRODUCED BY CD8+CCR5. ONE OF THE MAJOR HIV-SUPPRESSIVE HIV-1, HIV-2, AND SIMIAN IMMUNODEFICIENCY VIRUS (SIV).
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 Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.; "Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences."; Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
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TISSUE SPECIFICITY: T-CELL AND MACROPHAGE SPECIFIC.

INDUCTION: By mitogens.

SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY (SMALL
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MEDLINE=95244456; PubMed=7537088;
Skelton N.J., Aspiras F., Ogez J., Schall T.J.;
"Proton NMR assignments and solution conformation of RANTES, chemokine of the C-C type.";
Biochemistry 34:5329-5342(1995).
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MEDLINE=99111238; PubMed=9889151;
Wilken J., Hoover D., Thompson D.A., Barlow P.N., McSparron Picard L., Wlodawer A., Lubkowski J., Kent S.B.;
"Total chemical synthesis and high-resolution crystal struct the potent anti-HIV protein AOP-RANTES.";
Chem. Biol. 6:43-51(1999).
R.M.,
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MEDLINE=96106406; PubMed=8525373;
Cocchi F., DeVico A.L., Garzino-Demo A., Arya
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MEDLINE=95352612; PubMed=7542919;
Chung C.-W., Cooke R.M., Proudfoot A.E.I.,
"The three-dimensional solution structure
Biochemistry 34:9307-9314(1995).
                                                                                                                                                                                                                                                                                             Lusso P.;
"Identification of RANTES, MIP-1 alpha, a
HIV-suppressive factors produced by CD8+
Science 270:1811-1815(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          M21121; AAA36725.1; -.
AF043341; AAC03541.1; -.
AF088219; AAC63331.1; -.
AF266753; AAF73070.1; -.
BC008600; AAH08600.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC) .
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1HRJ; 14-OCT-96.

IRTN; 03-JUN-95.

1RTO; 03-JUN-95.

1B3A; 23-APR-99.
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EMBL;
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PDB;
PDB;
PDB;
PDB;
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MKVSAARLAVILIATALCAPASASPYSSDTTPCCFAYIARPL?RAHIKEYFYTSGKCSNP
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                                                                                                                                                                                                                                        L CYTOKINES CC; 1.
T-cell; Signal; Inflammatory response;
                              MIM; 18/011; -.

GO; GO:0003800; F:antiviral response protein activity; TAS.
GO; GO:0008009; F:chemokine activity; TAS.
GO; GO:0006874; P:calcium ion homeostasis; TAS.
GO; GO:0006928; P:cell adhesion; TAS.
GO; GO:0006928; P:cell motility; TAS.
GO; GO:000698; P:cellular defense response; TAS.
GO; GO:000698; P:chemotaxis; TAS.
GO; GO:0006987; P:exocytosis; TAS.
GO; GO:0006987; P:response to oxidative stress; TAS.
GO; GO:0006915; P:response to oxidative stress; TAS.
GO; GO:0009615; P:response to viruses; TAS.
GO; GO:0007165; P:signal transduction; TAS.
InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_ID8.
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                                                                                                                                                                                                                                                                                      SMALL INDUCIBLE CYTOKINE AS
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
Mammalia; Eutheria; Rodentia; Hystricognathi; Caviidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    databases
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Campbell E.M., Proudfoot A.E.I., Yoshimura T., Al
Wells T.N.C., White A.M., Westwick J., Watson M.L
Submitted (NOV-1996) to the EMBL/GenBank/DDBJ data
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P97272; 009076;
01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Small inducible cytokine A5 precursor (CCL5) (Tprotein) (SIS-delta).
CCL5 OR SCYAS.
CCL5 OR SCYAS.
CCL5 OR SCYAS.
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Pred. No. 7.3e-46;
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9990 MW;
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2; SMALL
1EQT; 19-AFK-00.
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E; PS00472; SMALI
ne; Chemotaxis; 7
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4 W 9 C 0 O
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404750
404760
60
                                                                                                                                                                                                                                                                3D-structure.
              Genew; HGNC:
MIM; 187011;
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DISULFID
CONFLICT
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Best Local (
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Matches
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10073 MW;
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                           (CHEMOKINE CC).
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                                                                                                                                                                                                                                                                                                                                                               91 AA;
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             SIMILARITY:
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                                                                                                                                                                                                                                                                                                                                                                                                                    73;
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Mammalia; E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOUSE
                                                                                                                                                                                                                                                                                        Cytokine;
                               ပ်
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61
                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                      SIGNAL
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                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                        Asano K., Nakamura M., Oguma T., Fukunaga K., Ishizaka A.,
Yamaguchi K., Kanazawa M.;
Submitted (APR-1997) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: CHEMOATTRACTANT FOR BLOOD MONOCYTES, MEMORY T HELPER
CELLS AND EOSINOPHILS. CAUSES THE RELEASE OF HISTAMINE FROM
BASOPHILS AND ACTIVATES EOSINOPHILS (BY SIMILARITY).
-!- SUBCELLULAR LOCATION: Secreted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (T-cell specific RANTES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                T HELPER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                   CYTOKINES CC; 1.
-cell; Signal; Inflammatory response POTENTIAL.
SMALL INDUCIBLE CYTOKINE A5.
BY SIMILARITY.
BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Intestine;
Aust G., Thamm B., Rost A.K.;
Cloning of Bos taurus RANTES mRNA.";
Submitted (MAR-1999) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: CHEMOATTRACTANT FOR BLOOD MONOCYTES, MEMORY T CELLS AND EOSINOPHILS. CAUSES THE RELEASE OF HISTAMINE BASOPHILS AND ACTIVATES EOSINOPHILS (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 91;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7F6A31B751237DB9 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 416; DB 1;
Pred. No. 1.3e-39;
2; Mismatches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Small inducible cytokine A5 precursor (CCLS)
protein) (SIS-delta).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; U77037; AAC53293.1; -.
EMBL; AB002662; BAA19604.1; -.
HSSP; P13501; 1RTN.
InterPro; IPR000827; CC_chemkine_sml
InterPro; IPR001811; Chemokine_ILB.
Pfam: PF00048; ILB; 1.
                                                                                                                                                                                                                                                                                                                                                            Interfic,
InterPro; IPROULGE.

Pfam; PF00048; IL8; 1.

SMART; SM00199; SCY; 1.

PROSITE; PS00472; SMALL_CYTOKINF
PROSITE; Chemotaxis; T-cell; S.

Trokine; Chemotaxis; T-cell; S.
SM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10088 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           87.0%;
larity 86.8%;
Conservative
                                                                                                                                                             (CHEMOKINE CC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 79; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EQUENCE FROM N.A.
              UENCE FROM N.A
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            SEQUENCE FRO TISSUE=Lung;
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097919;
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Local

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=92277990; PubMed=1375672;
Heeger P., Wolf G., Meyers C., Sun M.J., O'Farrell S.C.,
Krensky A.M., Neilson E.G.;
"Isolation and characterization of cDNA from renal tubular epithelium
encoding murine Rantes.";
Kidney Int. 41:220-225(1992).
LOCATION: Secreted.
BELONGS TO THE INTERCRINE BETA FAMILY (SMALL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (T-cell specific RANTES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                lus (mouse).
; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             S cytokine and man.";
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                                                                                                                                                                                                                                                                                                                                                                                                                  Inflammatory response
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MEDLINE=94132613; PubMed=7507961;
Danoff T.M., Lalley P.A., Chang Y.S., Heeger P.S., Neilson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 91;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         SMALL INDUCIBLE CYTOKINE AS BY SIMILARITY.
BY SIMILARITY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; DB 1; Le
2.7e-38;
9;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUL-1993 (Rel. 26, Created)
01-JUN-1994 (Rel. 29, Last sequence update)
15-SEP-2003 (Rel. 42, Last annotation update)
Small inducible cytokine A5 precursor (CCL5)
protein) (SIS-delta) (MuRantes).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       91
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  d. No. 2.76
Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                            KINES CC; 1.; Signal; In
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 404;
                                                                                                                                                                                                                                                                                                           sml.
                                                                                                                                                                                                                                   Chemotaxis; T-cell; Sign 1 23
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MEDLINE=92289805; PubMed=1376260;
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STRAIN-C57BL/6J; TISSUE-Pancreas;
Pubmed=1246881;
Pubmed=1246881;
Pubmed=1246881;
Pubmed=1246881;
Pubmed=1246881;
Pubmed=1246881;
Nikaido I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H., Yagi K., Tomaru Y., Hasegawa Y., Nogami A., Schonbach C., Golobori T., Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J., Schrinl L.M., Kanapin A., Matenda H., Batalov S., Beisel K.W., Black J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S., Dalla E., Dragani T.A., Fletcher C.F., Forrest A., Frazer K.S., Gasterland T., Cariboldi M., Gissi C., Godsin A., Guugh J., Grimmond S., Gustincich S., Hirokawa N., Jackson I.J., Jarvis E.D., Kanaja A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A., Maglott D.R., Maltais L., Marchionni L., McKenzie L., Miki H., Nagashima T., Numata K., Okido T., Pavan W.J., Pertea G., Pesole G., Petrovsky N., Pillai R., Pontius J.U., Qi D., Ramachandran S., Sandelin A., Schneider C., Semple C.A., Setou M., Shimada K., Sultana R., Takenaka Y., Taylor M.S., Tasafale R.D., Tomita M., Vang L., Yang L., Kawai J., Alzawa K., Arakawa T., Fukuda S., Hirozane-Kishikawa T., Konno H., Nakamura M., Sakazume N., Sato K., Shiraki T., Waki K., Kawai J., Alzawa K., Arakawa T., Fukuda S., Hara A., Hashizume W., Imotani K., Ishii Y., Itoh M., Kagawa I., Mayazaki A., Sasaki D., Shibata K., Shinagawa A., Sasaki D., Shibata K., Shinagawa A., Sasaki D., Shibata K., Shinagawa A., Sasaki D., Shinagawa A., Sasaulahi D., Shibata K., Shinagawa A., Sasaki D., Shibata W., Sakazuma M., Sasaki D., Shibata W., Sakazuma M., Sasaki D., Shibata W., Sakazuma M., Sakazuma M., Sakazuma M., Sakazuma M., Sakazuma M., Saka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Sequence polymorphisms in the chemokines Scyal (TCA-3), Scya2 (monocyte chemoattractant protein (MCP)-1), and Scyal2 (MCP-5) are candidates for eae7, a locus controlling susceptibility to monophasic remitting/nonrelapsing experimental allergic encephalomyelitis.";
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localization
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Nomiyama H.;
"Organization of the mouse CC chemokine cluster containi for Cl0, MRP-2 and RANTES.";
Submitted (NOV-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                          PubMed=7513046;
e B.E., Shin M.L., Noble P.W., Fisher
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F.,
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and chromosomal lo chemokine RANTES.";
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STRAIN=B10.S/J, BALB/CJ, DBA/2J, NOD/LTJ, and
MEDLINE=99370037; PubMed=10438970;
Teuscher C., Butterfield R.J., Ma R.Z., Zachar
  encoding the murine (152:1182-1189(1994))
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"Analysis of the mouse trans
60,770 full-length cDNAs.";
Nature 420:563-573(2002).
                                                                                                                                                                                                                                      Drysdale
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MEDLINE=94217689;
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Blankenhorn E.P.;
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Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J., Bosak S.A., McEwan P.J., McKernen K.J., Malek J.A., Gunaratne P.H., A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., A Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A., Shoting A.C., Shovchenko Y., Bouffard G.G., A Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C., A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E., A Schnerch A., Schein J.E., Jones S.J.M., Marra M.A., Inflance C.D., Dickson M.C., Touchman initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";

Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).

-!- FUNCTION: CHEMOATTRACTANT FOR BLOOD MONOCYTES, MEMORY T HELPER CELLS AND BOSINOPHILS. CAUSES THE RELEASE OF HISTAMINE FROM BASOPHILS AND ACTIVATES EOSINOPHILS.

-!- FUNCTION: Secreted.

-!- TISSUE SPECIFICITY: T-CELL AND MACROPHAGE SPECIFIC.

-!- SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY (SMALL CYTOKINE C.C.) (CHEMORINE CC).
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cis; T-cell; Signal; Inflammatory response
23 POTENTIAL.
91 SMALL INDUCIBLE CYTOKINE A5.
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SDFD66F4684FE1C8 CRC64;
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EMBL; S37648; AAA18302.1; -.
EMBL; W70675; CAA50011.1; -.
EMBL; AF065944; AAC17511.1; -.
EMBL; AF065944; AAC17512.1; -.
EMBL; AF065944; AAC17512.1; -.
EMBL; AF065947; AAC17512.1; -.
EMBL; AF065947; AAC17512.1; -.
EMBL; AF065947; AAC17513.1; -.
EMBL; AF065947; AAC17513.1; -.
EMBL; AF065947; AAC17514.1; -.
EMBL; AF065947; AAC17514.1; -.
EMBL; AF065947; AAF25528.1; -.
EMBL; AF065947; AAF2558.1; -.
EMBL; AF0633508; AAH33508.1; -.
EMBL; AF0033508; AAH33508.1; -.
EMBL; AF0033508; AAH33508.1; -.
EMBL; AF000927; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_IL8.
InterPro; IPR001811; Chemokine_IL8.
InterPro; IPR001811; Chemokine_IL8.
Cytokine; Chemotaxis; T-cell; Signal;
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79.1%;
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                                                                                                                                                                                                                  STRAIN=Long Evans; TISSUE=Lung;
Jones M.L., Shanley T.P., Ward P.A.;
Submitted (FEB-1994) to the EMBL/GenBank/DDBJ databases.

-!- FUNCTION: CHEMOATTRACTANT FOR BLOOD MONOCYTES, MEMORY T HELPER
CELLS AND EOSINOPHILS. CAUSES THE RELEASE OF HISTAMINE FROM
BASOPHILS AND ACTIVATES EOSINOPHILS (BY SIMILARITY).

-!- SUBCELLULAR LOCATION: Secreted.

-!- SUBCELLULAR LOCATION: Secreted.

-!- SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY (SMALL CYTOKINE C-C) (CHEMOKINE CC).
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igmodon hispidus (Hispid cotton rat).
ukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi
lammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Sigmodontin
                                                                                                  Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Rai
NCBI_TaxID=10116,
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                                                       (T-cell
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InterPro; IPR001811; Chemokine_IL8.
Pfam; PF00048; IL8; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.
Cytokine; Chemotaxis; T-cell; Signal; Inflammatory SIGNAL
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Pred. No. 5.3e-36;
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28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Small inducible cytokine A5 precursor (CCL5)
protein).
01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Small inducible cytokine A5 precursor (CCL5)
protein) (SIS-delta).
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ilarity 78.3%;
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HSSP; P13501; 1RTN.
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72; Conser
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01-NOV-1995 (Rel. 32, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Small inducible cytokine A4 precursor (CCL4) (Macrophage inflammatory protein 1-beta) (MIP-1-beta) (Immune activation protein 2) (ACT-2).
                                                                                                                                                                                                                                                            (SMALL CYTOKINE
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STRAIN=New Zealand white;

MEDLINE=94198229; PubMed=8148323;

Mori S., Goto K., Goto F., Mutakami K., Ohkawara S., Yoshinaga M.;

"Dynamic changes in mRNA expression of neutrophils during the course of acute inflammation in rabbits.";

Int. Immunol. 6:149-156(1994).

-!- FUNCTION: MONOKINE WITH INFLAMMATORY AND CHEMOKINETIC PROPERTIES (BY SIMILARITY).
                                                                   Blanco J.C., Pletneva L.M., Prince G.A.;
"Sigmodon hispidus cytokines, chemokines and interferons.";
Submitted (SEP-2001) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: CHEMOATTRACTANT FOR BLOOD MONOCYTES, MEMORY T HELPER
CELLS AND EOSINOPHILS. CAUSES THE RELEASE OF HISTAMINE FROM
BASOPHILS AND ACTIVATES EOSINOPHILS (BY SIMILARITY).
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY (SMALL CYTOK).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
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Pfam; PF00048; ILB; 1.
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PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.
PROSITE; PS00472; SMALL_CYTOKINE A5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sml.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; AF421391; AAL16932.1; -.
InterPro; IPR000827; CC_chemkine_si
InterPro; IPR001811; Chemokine_IL8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11;
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                                                                                                                                                                                                                                                                                      (CHEMOKINE CC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                  FROM N.A
TaxID=42415;
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                                                SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SMALL CYTOKINE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              genes are closely linked and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein) (PAT 464.2).
CCL3L1 OR SCYA3L1 OR GOS19-2.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=90287702; PubMed=1972563; Irving S.G., Zipfel P.F., Balke J., McBride O.W., Morton Burd P.R., Siebenlist U., Kelly K.; "Two inflammatory mediator cytokine genes are closely linvariably amplified on chromosome 17q."; Nucleic Acids Res. 18:3261-3270(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                   Inflammatory response; Signal.
Inflammatory response; Signal.
BY SIMILARITY.
SMALL INDUCIBLE CYTOKINE
BY SIMILARITY.
BY SIMILARITY.
                        FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ECBA8818D42A735C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ore 254.5; DB 1; ed. No. 1.2e-21; Mismatches 28;
                      BETA
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MEDLINE=91103879; PubMed=2271120;
Blum S., Forsdyke R.E., Forsdyke D.R.;
"Three human homologs of a murine gene
                                                                                                                                                                                                                                                                    EMBL; D17402; BAA04226.1; -.
PIR; I46730; I46730.
HSSP; P13236; IHUM.
InterPro; IPR000827; CC_chemkine_sml
InterPro; IPR001811; Chemokine_IL8.
Pfam; PF00048; IL8; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL CYTOKINES_CC
Cytokine; Chemotaxis; Inflammatory r
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score
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Cell Biol. 9:589-602(1990)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       53.2%;
illarity 51.1%;
Conservative 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10066 MW;
SUBCELLULAR LOCATION: SIMILARITY: BELONGS TO
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                                                (CHEMOKINE
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TISSUE=Blood;
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SIGNAL
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DISULFID
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                                                                                                                                                          GO; GO: 0005576; C: extracellular; NAS.
GO; GO: 0008009; F: chemokine activity; NAS.
GO; GO: 0006935; P: chemotaxis; NAS.
GO; GO: 0006954; P: inflammatory response; TAS.
GO; GO: 0008285; P: negative regulation of cell proliferation; TAS.
InterPro; IPR000827; CC chemkine sml.
InterPro; IPR001811; Chemokine_IL8.
Pfam; PF00048; IL8; 1.
SMART; SM00199; SCY; 1.
SMART; SM00199; SCY; 1.
Cytokine; Chemotaxis; Signal.
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                                               cytokine LD78 and their
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A7A79E774006D61E CRC64;
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Pred. No. 2.1e-21;
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MEDLINE=90287155; PubMed=1694014;
Nakao M., Nomiyama H., Shimada K.;
"Structures of human genes coding for
expression.";
                                                                                                   Cell. Biol. 10:3646-3658(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; X52149; CAA36397.1; -.
EMBL; M24110; AAA35859.1; -.
EMBL; D90145; BAA14173.1; -.
EMBL; BC007783; AAH07783.1; -.
EMBL; BC027888; AAH27888.1; -.
PIR; B35673; B35673.
HSSP; P13236; 1HUM.
Genew; HGNC:10628; CCL3L1.
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MIM, 601395;
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Best Local
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1 MKVSAARLAVILIATALCAPASASPYSSDT-TPCCFAYIARPLPRAHIKEYFYTSGKCSN

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Matches
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                                                                                                                                                                                   inflammatory
                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
TISSUE=Bone marrow;
MEDLINE=95369710; PubMed=7642115;
Petrenko O., Ischenko I., Enrietto P.J.;
"Isolation of a cDNA encoding a novel chicken chemokine homologous to mammalian macrophage inflammatory protein-1 beta.";
Gene 160:305-306(1995).
 PROPERTIES
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                                                                                                                             SY04 CHICK STANDARD; PRT; 90 AA;
SY04 CHICK STANDARD; PRT; 90 AA;
Sy0826; O910C9;
O1-NOV-1997 (Rel. 35, Created)
28-FEB-2003 (Rel. 41, Last annotation update)
28-FEB-2003 (Rel. 41, Last annotation update)
Small inducible cytokine A4 homolog precursor (Macrophage inflammator protein 1-beta homolog).
CCL4 OR SCYA4.
Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
Gallus.
                                                                                                                                                                                                                                                                                                                                                                                                                   the mammalian
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INFLAMMATORY AND CHEMOKINETIC
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SEQUENCE OF 14-90 FROM N.A.
Petrenko O., Enrietto P.J.;
Submitted (JUL-1994) to the EMBL/GenBank/DDBJ database
-!- FUNCTION: MONOKINE WITH INFLAMMATORY AND CHEMOKINE
-!- SUBUNIT: Homodimer (By similarity).
-!- SUBCELLULAR LOCATION: Secreted.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CRC64;
                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
Hughes S.M., Bumstead N.;
"Mapping of the gene encoding the chicken homologue chemokine SCYA4.";
Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databa
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SMALL INDUCIBLE (
BY SIMILARITY.
BY SIMILARITY.
M -> L (IN REF. 1
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                                                     PAVVFVTRKNRQVCANPEKKWVREYINSLEMS
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EMBL; AJ243034; CAB45103.1; -.
HSSP; P13236; 1HUM.
InterPro; IPR000827; CC_chemkine_sml
InterPro; IPR001811; Chemokine_IL8.
Pfam; PF00048; IL8; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC
Cytokine; Chemotaxis; Signal.
SIGNAL
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01-MAR-1989 (Rel. 10, Created)
01-MAR-1989 (Rel. 10, Last sequence update)
15-SEP-2003 (Rel. 42, Last annotation update)
15-SEP-2003 (Rel. 42, Last annotation update)
Small inducible cytokine A3 precursor (CCL3) (Macrophage inflammatory Small inducible cytokine A3 precursor (CCL3) (Macrophage inflammatory protein 1-alpha) (MIP-1-alpha) (Tonsillar lymphocyte LD78 alpha protein 1-alpha) (MIP-1-alpha) (Tonsillar lymphocyte LD78 alpha protein) (G0/G1 switch regulatory protein 19-1) (G0S19-1 protein)
(SIS-beta) (PAT 464.1).
(CL3 OR SCYA3 OR G0S19-1 OR MIP1A.
Homo sapiens (Human).
                                                                    MKVSAARLAVILIATALCAPASASPYSSD-TTPCCFAYIARPLPRAHIKEYFYTSGKCSN
                                                                                                       stem
                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=89140347; PubMed=2521882; Zipfel P.F., Siebenlist U.; Zipfel P.F., Balke J., Irving S.G., Kelly K., Siebenlist U.; "Mitogenic activation of human T cells induces two closely related genes which share structural similarities with a new family of secreted factors.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=91103879; PubMed=2271120;
Blum S., Forsdyke R.E., Forsdyke D.R.;
"Three human homologs of a murine gene encoding an inhibitor of
cell proliferation.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=96127782; PubMed=8541527;
Hunter M.G., Bawden L., Brotherton D., Craig S., Cribbes S., Czaplewski L.G., Dexter T.M., Drummond A.H., Gearing A.H., Heyworth C.M., Lord B.I., Mccourt M., Varley P.G., Wood L.M., Edwards R.M., Lewis P.J.;
"BB-10010: an active variant of human macrophage inflammatory 1 alpha with improved pharmaceutical properties.";
Blood 86:4400-4408(1995).
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MEDLINE=86223879; PubMed=3086300;
Obaru K., Fukuda M., Maeda S., Shimada K.;
Obaru K., Fukuda M., Maeda S., Shimada K.;
"A cDNA clone used to study mRNA inducible in human tonsillar
"A cDNA clone used to study mRNA inducible in human tonsillar
lymphocytes by a tumor promoter.";
Jymphocytes by a tumor promoter.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases
                         Indels
Pred. No. 2.6e-21; ; Mismatches 23;
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                                                                                                                                                                                             60 PAVVFVTRKNRQVCANPEKKWVREYINSLEMS
                                                                                                                                                                                                                           AGVVFITRKGREVCANPENDWVQDYMNKMELN
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Mol. Cell. Biol. 10:3646-3658(1990).
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"Structures of human genes coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE=90287155; PubMed=1694014;
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 51.1%;
                                                                                                                                                                                                                                                                                                                                                                                            STANDARD;
                              Conservative
      Local Similarity
les 47; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9606;
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MITOGEN)).

-!- SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY (SMALL CYTOKINE C-C) (CHEMOKINE CC). STRONG, TO SCYA3L1.

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                                                 "Identification of RANTES, MIP-1 alpha, and MIP-1 beta as the major HIV-suppressive factors produced by CD8+ T cells.";
Science 270:1811-1815(1995).
-!- FUNCTION: MONOKINE WITH INFLAMMATORY AND CHEMOKINETIC PROPERTIES.
BINDS TO CCR1, CCR4 AND CCR5. ONE OF THE MAJOR HIV-SUPPRESSIVE FACTORS PRODUCED BY CD8+ T CELLS. RECOMBINANT MIP-1-ALPHA INDUCES A DOSE-DEPENDENT INHIBITION OF DIFFERENT STRAINS OF HIV-1, HIV-2, AND SIMIAN IMMUNODEFICIENCY VIRUS (SIV).
-!- SUBCELLULAR LOCATION: Secreted.
-!- INDUCTION: BY TPA OR PHA (TPA = 12-O-TETRADECANOYL PHORBOL-13 ACETATE (TUMOR PROMOTER); PHA = PHYTOHEMAGGLUTININ (T-CELL MITOGEN)).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        response; Signal; 3D-structure.
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EMBL; M234525.1 AA31616.1; --
EMBL; M234525.1 AA31616.1; --
EMBL; M23452.2 AA31616.1; --
EMBL; M23452.2 AA31616.1; --
EMBL; M20118; AA47255.1; --
EMBL; M20118; AA47255.1; --
EMBL; M20118; AA47255.1; --
EMBL; M20118; AA44172.1; --
EMBL; M23178; AA44172.1; --
EMBL; M23573; A30574.
PDB; 1B53; 22-JUL-99.
PDB; 1B54; 2A50, 2A50,
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BY SIMILARITY.
BY SIMILARITY.
D->A: IN BB-10010; IMPROVED PHARMACEUTICAL PROPERTIES.
; 517865D5D6776CA8 CRC64;
                                   Gallo
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            d=8525373;
Garzino-Demo A., Arya S.K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 248; DB 1; I
Pred. No. 6.5e-21;
22; Mismatches 25;
  PubMed=852537
A.L., Garzino
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imilarity 48.4%;
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SCY; 1.
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MEDLINE=96106406;
Cocchi F., Devico
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SEQUENCE OF 24-5).

MEDLINE=96183056; PubMed=8607872;

MEDLINE=10duced neutrophil chemoattractant (CINC)-2 alpha, a novel

"Cytokine-induced by a predominant chemokine produced by

"Commun 220:945-948(1996).

"Ipopolysaccharide-stimulated rat macrophages in culture.";

Biochem. Biophys. Res. Commun. 220:945-948(1996).

"Indepolysaccharide-stimulated rat macrophages in culture.";

"Commun 220:945-948(1996).

"Comm
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MKVSAARLAVILIATALCAPASASPYSSDTTPCCFAYIARPLPRAHIKEYFYTSGKCSNP
                                   (Macrophage inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PROPERTIES
EOSINOPHILS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE=Lung;
MEDLINE=95298037; PubMed=7779098;
Shi M.M., Godleski J.J., Paulauskis J.D.;
Shi M.M., Godleski J.J., Paulauskis J.D.;
"Molecular cloning and posttranscriptional regulation of macrophage "Molecular cloning and posttranscriptional regulation of macrophage inflammatory protein-1 alpha in alveolar macrophages.";
Inflammatory protein-1 alpha in alveolar macrophages.";
Riochem. Biophys. Res. Commun. 211:289-295(1995).
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1 alpha) in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
STRAIN=Long Evans; TISSUE=Lung;
MEDLINE=95238980; PubMed=7722328;
Shanley T.P., Schmal H., Friedl H.P., Jones M.L., Ward F
"Role of macrophage inflammatory protein-1 alpha (MIP-1
                                                                                                                                                                                                                                                                                                                                                                          01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Small inducible cytokine A3 precursor (CCL3)
protein 1-alpha) (MIP-1-alpha).
CCL3 OR SCYA3 OR MIP1A.
Rattus norvegicus (Rat).
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INDUCTION: BY LIPOPOLYSACCHARIDE (LPS)
SIMILARITY: BELONGS TO THE INTERCRINE
C-C) (CHEMOKINE CC).
                                                                                                                                                    |:|:|:::|||||:| ::||::||:| GVIFLTKRSRQVCADPSEEWVQKYVSDLELS
                                                                                                                        AVVFVTRKNRQVCANPEKKWVREYINSLEMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   te lung injury in rats.";
Immunol. 154:4793-4802(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; U22414; AAA80608.1; -. EMBL; U06435; AAA96498.1; -. PIR; I52322; I52322. HSSP; P13236; 1HUM.
                                                                                                                                                                                                                                                                                                                               STANDARD;
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"A family of small inducible proteins secreted by leukocytes are members of a new superfamily that includes leukocyte and fibroblast-derived inflammatory agents, growth factors, and indicators of various activation processes.";
J. Immunol. 142:679-687(1989).
                                                                                                                                                                       SN
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                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=89067830; PubMed=3058856; Sherry B., Tekamp-Olson P., Gallegos C., Bauer D., Davatelis G., Sherry B., Tekamp-Olson P., Gallegos C., Bauer D., Davatelis G., Wolpe S.D., Masiarz F., Coit D., Cerami A.; "Resolution of the two components of macrophage inflammatory protein 1, and cloning and characterization of one of those components, macrophage inflammatory protein 1 beta."; J. Exp. Med. 168:2251-2259(1988).
                                                                                                                                                    Gaps
                                                                                                                                                                                inflammator (ACT2).
                                               Signal; Heparin-binding
                                                                                                                                                                       MKVSAARLAVILIATALCAPASASPYSSDT-TPCCFAYIARPLPRAHIKEYFYTSGKC
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: Murinae; Mu
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Scya12 (MCP-5)
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                                                                                                                                                                                                                                                                                                                        (Macrophage (SIS-gamma)
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                                                                                                                                  Length
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                                                                                                    AND 3)
CRC64;
                                                                  CYTOKINE
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                                                                                                                           ; DB 1; Le
6.5e-21;
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                                                                                           -> T (IN REF. 2)
-> W (IN REF. 2 )
14E861C647F9A2EB (
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                                                                                                                                                                                                                                                                                          01-JAN-1990 (Rel. 13, Created)
01-FEB-1996 (Rel. 33, Last sequence update)
15-SEP-2003 (Rel. 42, Last annotation update)
Small inducible cytokine A4 precursor (CCL4)
protein 1-beta) (MIP-1-beta) (H400 protein) (CCL4 OR SCYA4 OR MIP1B.
Mus musculus (Mouse).
                                                                  INDUCIBLE
                                               response;
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                                                                                                                                                                                                                            9
                                                                 SMALL INDUCIBL BY SIMILARITY. BY SIMILARITY.
                                                                                                                                                    Mismatches
                                                                                                                                                                                                           PAVVFVTRKNROVCANPEKKWVREYINSLEMS
                                                                                                                                                                                                                     .
В.
                                                                                                                                  Score 248;
Pred. No.
le sml.
IL8.
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                                       ບິ
                                      S00472; SMALL_CYTOKINES_(Chemotaxis; Inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
MEDLINE=89093958; PubMed=2521353;
 CC_chemkine
Chemokine_]
                                                                                                                                                  18;
                                                                                                                X.
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Daubersies P., Lepretre F.,
                                                                                                                                51.9%;
51.1%;
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(OCT-1991) to
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SCY; 1.
SMALL
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4828
7687
                ; ;
InterPro; IPR000827; (InterPro; IPR001811; OPfam; PF00048; IL8; ISSMART; SM00199; SCY; IPROSITE; PS00472; SMAI
                                                                                                                                                                                                                                                                                                                                                           Metazoa;
Eutheria;
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NCBI TaxID=1
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                                              Cytokine;
SIGNAL
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                     SUBCELLULAR LOCATION: Secreted.
SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY (SMALL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (Macrophage inflammatory pha) (Heparin-binding
candidates for eae7, a locus controlling susceptibility to monophasic remitting/nonrelapsing experimental allergic encephalomyelitis.";
J. Immunol. 163:2262-2266(1999).
-!- FUNCTION: MONOKINE WITH INFLAMMATORY AND CHEMOKINETIC PROPERTIES.
-!- SUBUNIT: Homodimer (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Euteleostomi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleosto
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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01-JUL-1989 (Rel. 11, Created)
01-APR-1990 (Rel. 14, Last sequence update)
15-SEP-2003 (Rel. 42, Last annotation update)
Small inducible cytokine A3 precursor (CCL3) (Macriprotein 1-alpha) (MIP-1-alpha) (TY-5) (SIS-alpha)
chemotaxis protein) (L2G25B).
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Pred. No. 1.2e-20;
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HSSP; P13236; 1HUM.
MGD; MGI:98261; Cc14.
InterPro; IPR000827; CC_chemkine_sml
InterPro; IPR001811; Chemokine_IL8.
InterPro; IPR001811; Chemokine_IL8.
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M00199; SCY; 1.

PS00472; SMALL CYTOKINES CC;

; Chemotaxis; Inflammatory r

1 23 SMALL I

24 92 SMALL I

34 58 BY SIMI

34 58 BY SIMI

35 74 BY SIMI
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EMBL; M35590; AAA39708.1; -.
EMBL; X62502; CAA44364.1; -.
EMBL; AF128218; AAF22559.1; -.
EMBL; AF128219; AAF22560.1; -.
PIR; C30552; C30552.
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SMART; SM00199; SCY,
PROSITE; PS00472; SN
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STRAIN=BALB/CJ, DBA/2J, NOD/LTJ, SJL/J, and B10.S/J; TISSUE=Spleen;
MEDLINE=99370037; PubMed=10438970;
Teuscher C., Butterfield R.J., Ma R.Z., Zachary J.F., Doerge R.W.,
Blankenhorn E.P.;
"Sequence polymorphisms in the chemokines Scyal (TCA-3), Scya2
(monocyte chemoattractant protein (MCP)-1), and Scyal2 (MCP-5) are
candidates for eae7, a locus controlling susceptibility to monophasi
remitting/nonrelapsing experimental allergic encephalomyelitis.";
J. Immunol. 163:2262-2266(1999).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           alpha
                                                                                                                                                                                                                                                                                                                            MEDLINE=91016858; PubMed=2216738; Grove M., Lowe S., Graham G., Pragnell I., Plumb M.; Grove M., Lowe S., Graham G., Pragnell I., Plumb M.; "Sequence of the murine haemopoietic stem cell inhibitor/macrophage inflammatory protein 1 alpha gene."; Nucleic Acids Res. 18:5561-5561(1990).
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FOR EOSINOPHILS
CALCIUM RELEASE
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TISSUE SPECIFICITY: EXPRESSED IN LUNG, SPLEEN, AND PANCREAS SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY (SMALL CY. C-C) (CHEMOKINE CC).
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                                                      Luedke
                                                                                                                        SEQUENCE FROM N.A.
MEDLINE=89093958; PubMed=2521353;
Brown K.D., Zurawski S.M., Mosmann T.R., Zurawski G.;
"A family of small inducible proteins secreted by leukocytes members of a new superfamily that includes leukocyte and fibroblast-derived inflammatory agents, growth factors, and indicators of various activation processes.";
J. Immunol. 142:679-687(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=88154745; PubMed=3279154;
MeDLINE=88154745; PubMed=3279154;
Wolpe S.D., Davatelis G., Sherry B., Beutler B., Hesse D.G.,
Nguyen H.T., Moldawer L.L., Nathan C.F., Lowry S.F., Cerami
"Macrophages secrete a novel heparin-binding protein with
inflammatory and neutrophil chemokinetic properties.";
J. Exp. Med. 167:570-581(1988).
-!- FUNCTION: MONOKINE WITH INFLAMMATORY, PYROGENIC AND CHEM
PROPERTIES. HAS A POTENT CHEMOTACTIC ACTIVITY FOR EOSINC
BINDING TO A HIGH-AFFINITY RECEPTOR ACTIVATES CALCIUM RE
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MEDLINE=91237116; PubMed=2033269;
Widmer U., Yang Z., van Deventer S., Manogue K.R.,
Cerami A.;
"Genomic structure of murine macrophage inflammaton
                                                   .D., Hermser
Cerami A.;
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SEQUENCE FROM N.A.
MEDLINE=89184547; PubMed=2784565;
Kwon B.S., Weissman S.M.;
Kwon B.S., Weissman S.M.;
"cDNA sequences of two inducible T-cell genes.";
"cDNA sequences of two inducible T-cell genes.";
                                                      S,D
                                               Wolpe S.
       Colons.

REVISIONS.
Davatelis G., Tekamp-Olson P., Wolp Gallegos C., Coit D., Merryweather Med. 170:2189-2189(1989).
  167:1939-1944(1988)
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Y04 HUMAN STANDALL.

C P13236; P22617; Q13704;

C P13236; P22617; Q13704;

T 01-JAN-1990 (Rel. 13, Last sequence update)

Y1 01-JAN-1990 (Rel. 13, Last annotation update)

Y2 01-JAN-1990 (Rel. 13, Last annotation update)

Y3 15-SEP-2003 (Rel. 42, Last annotation update)

Y4 15-SEP-2003 (Rel. 42, Last annotation update)

Y5 Small inducible cytokine A4 precursor (CCL4) (Macrophage inflammatory

DE protein 1-beta) (MIP-1-beta) (T-cell activation protein 2) (ACT-2)

DE PAT 744) (H400) (SIS-gamma) (Lymphocyte activation gene-1 protein)

CCL4 OR SCYA4 OR MIP1B OR LAGI.

CCL4 OR SCYA4 OR MIP1B OR LAGI.

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MEDLINE=89140347; PubMed=2521882;
Zipfel P.F., Balke J., Irving S.G., Kelly K., Siebenlist U.;
Zipfel P.F., Balke J., Irving S.G., Kelly K., Siebenlist U.;
"Mitogenic activation of human T cells induces two closely related genes which share structural similarities with a new family of secreted factors.";
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SEQUENCE FROM N.A.
MEDLINE=89071764; PubMed=2462251;
Lipes M.A., Napolitano M., Jeang K.-T., Chang N.T.,
Lipes M.A., Napolitano M., Jeang K.-T., Chang N.T.,
"Identification, cloning, and characterization of a:
"Identification, cloning, and characterization of a:
"Identification, cloning, and characterization of a:
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Pred. No. 2.3e-20;
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PGVIFLTKRNRQICADSKETWVQEYITDLELN 91
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EMBL; X12531; CAA31047.1; -.
EMBL; X12531; CAA37452.1; -.
EMBL; X53372; CAA37452.1; -.
EMBL; X53372; CAA37452.1; -.
EMBL; AF065939; AAC17506.1; -.
EMBL; AF065940; AAC17508.1; -.
EMBL; AF065941; AAC17508.1; -.
EMBL; AF065941; AAC17508.1; -.
EMBL; AF065942; AAC17508.1; -.
EMBL; AF065943; AAC17509.1; -.
EMBL; AF065943; AAC17509.1; -.
EMBL; AF065943; AAC17509.1; -.
EMBL; AF065943; AC17509.1; -.
EMBL; AF065943; AAC17500.1; -.
EMBL; AF065941; AAC17500.1; -.
EMBL; AF065940; AAC17500.1; -.
EMBL; AF065941; AAC17500.1; -.
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InterPro; IPRUVICE
Pfam; PF00048; ILB; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL CYTOKINES CC;
PROSITE; Chemotaxis; Inflammatory reconnections and the second states of th
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10345 MW;
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142:1582-1590(1989)

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SEQUENCE FROM N.A.
MEDLINE=89093958; PubMed=2521353;
Brown K.D., Zurawski S.M., Mosmann T.R., Zurawski G.;
Brown K.D., Zurawski S.M., Mosmann T.R., Zurawski G.;
"A family of small inducible proteins secreted by leukocytes are members of a new superfamily that includes leukocyte and fibroblast-derived inflammatory agents, growth factors, and indicators of various activation processes.";
J. Immunol. 142:679-687(1989).
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STRUCTURE BY NMR.
MEDLINE=94182137; PubMed=8134838;
Lodi P.J., Garrett D.S., Kuscewski J., Tsang M.L.S., Weatherbee J.A., Lodi P.J., Garrett D.S., Kuscewski J., Tsang M.L.S., Weatherbee J.A., Lodi P.J., Garrett D.S., Kuscewski J., Tsang M.L.S., Weatherbee J.A., Lodia P.J., Garrett D.S., Kuscewski J., Tlore G.M.;
Leonard W.J., Gronenborn A.M., Clore G.M.;
"High-resolution solution structure of the beta chemokine hMIP-1 beta by multidimensional NMR.";

Science 263:1762-1767(1994).
Science 263:1762-1767(1994).
-!- FUNCTION: MONOKINE WITH INFLAMMATORY AND CHEMOKINETIC PROPERTIES.
BINDS TO CCRS AND TO CCR8. ONE OF THE MAJOR HIV-SUPPRESSIVE
BINDS TO CCR5 AND TO CCR8. ONE OF THE MAJOR HIV-SUPPRESSIVE

BINDS TO CCR5 AND TO CCR8. ONE OF THE MAJOR HIV-SUPPRESSIVE
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MEDLINE=98180363; PubMed=9521068;

Bernardini G., Hedrick J., Sozzani S., Luini W., Spinetti G.,

Weiss M., Menon S., Zlotnik A., Mantovani A., Santoni A.,

Napolitano M.;

"Identification of the CC chemokines TARC and macrophage inflammatory
protein-1 beta as novel functional ligands for the CCR8 receptor.";
                                                                                                                                                                                                                                                                                                                                                                                                       Genomic structure, HTLV-I/Tax
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lusso P.;
"Identification of RANTES, MIP-1 alpha, and MIP-1 beta as the major
HIV-suppressive factors produced by CD8+ T cells.";
Science 270:1811-1815(1995).
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                                                                                                                                  SEQUENCE FROM N.A.
MEDLINE=91061800; PubMed=2247088;
Baixeras E., Roman-Roman S., Jitsukawa S., Genevee C., Mechiche S., Viegas-Pequignot E., Hercend T., Triebel F.;
Viegas-Pequignot E., Hercend T., Triebel F.;
"Cloning and expression of a lymphocyte activation gene (LAG-1).";
Mol. Immunol. 27:1091-1102(1990).
                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
MEDLINE=91373378; PubMed=1894635;
Napolitano M., Modi W.S., Cevario S.J., Gnarra J.R., Seuanez H.N.,
Leonard W.J.;
"The gene encoding the Act-2 cytokine. Genomic structure, HTLV-I/Ti
responsiveness of 5' upstream sequences, and chromosomal
localization.";
J. Biol. Chem. 266:17531-17536(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE OF 6-92 FROM N.A.
MEDLINE=90038522; PubMed=2809212;
Miller M.D., Hata S., Waal Malefyt R., Krangel M.S.;
"A novel polypeptide secreted by activated human T lymphocytes.";
J. Immunol. 143:2907-2916(1989).
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structure on
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MEDLINE=96106406; PubMed=8525373;
Cocchi F., DeVico A.L., Garzino-Demo A., Arya S.K., Gallo R.C.,
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TISSUE=T-cell;
MEDLINE=89325421; PubMed=2568930;
Chang H.C., Reinherz E.L.;
"Isolation and characterization of a cDNA encoding a cytokine which is induced by stimulation via the CD2 human T lymphocytes.";
Eur. J. Immunol. 19:1045-1051(1989).
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Fasman K., McKernan K., Nusbaum C.,
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Birren B., Fasn
Lander E.;
Submitted (JAN-
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                                                                                       (SMALL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Signal; 3D-structure
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GO, GO: 0008009; F: chemokine activity; TAS.
GO; GO: 0004716; F: receptor signaling protein tyrosine kinase...
GO; GO: 0008151; P: cell adhesion; TAS.
GO; GO: 0008151; P: cell adventh and/or maintenance; TAS.
GO; GO: 0007267; P: cell motility; TAS.
GO; GO: 0007267; P: cell-cell signaling; TAS.
GO; GO: 0007267; P: immune response; TAS.
GO; GO: 0006955; P: immune response; TAS.
GO; GO: 0006955; P: inflammatory response; TAS.
GO; GO: 0006915; P: response to viruses; TAS.
GO; GO: 0008166; P: viral replication; TAS.
GO; GO: 0008166; P: viral replication; TAS.
InterPro; IPR001811; Chemokine_IL8.
InterPro; IPR001811; Chemokine_IL8.
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М
HIV-1,
                                                                       INDUCTION: By mitogens.
SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY
C-C) (CHEMOKINE CC).
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Inflammatory response;
DOSE-DEPENDENT INHIBITION OF DIFFERENT AND SIMIAN IMMUNODEFICIENCY VIRUS (SIV) SUBUNIT: Homodimer.
SUBCELLULAR LOCATION: Secreted.
                                                                                                                                                                                                                                                                              EMBL; M23502; AAA36656.1; -.
EMBL; M25316; AAA57256.1; -.
EMBL; J04130; AAA51576.1; -.
EMBL; X53683; CAA37723.1; -.
EMBL; X53682; CAA37722.2; ALT_SEQ.
EMBL; X16166; CAA34291.1; -.
EMBL; M69203; AAB00790.1; -.
EMBL; M69201; AAB00790.1; JOINED.
EMBL; M69202; AAB00790.1; JOINED.
EMBL; M69202; AAB00790.1; JOINED.
EMBL; M69202; AAB00790.1; JOINED.
EMBL; M57503; AAA36752.1; -.
PDB; 1HUM; 30-APR-94.
PDB; 1HUM; 30-APR-94.
PDB; 1JE4; 03-OCT-01.
Genew; HGNC:10630; CCL4.
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STRAIN=Long Evans; TISSUE=Lung;
Jones M.L., Shanley T.P., Schmal H., Friedl H.P., Ward P.A.;
Jones M.L., Shanley T.P., Schmal H., Friedl H.P., Ward P.A.;
Submitted (FEB-1994) to the EMBL/GenBank/DDBJ databases.

-!- FUNCTION: MONOKINE WITH INFLAMMATORY AND CHEMOKINETIC PROPERTIES

-!- SUBUNIT: Homodimer (By similarity).

-!- SUBCELLULAR LOCATION: Secreted.

-!- SUBCELLULAR LOCATION: Secreted.

-!- SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY (SMALL CYTOKIN)

C-C) (CHEMOKINE CC).
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Murinae; Rattus
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# HSSP; P13236; 1HUM.

# RSP; P13236; 1HUM.

# InterPro; IPR000827; CC_chemkine_sml.

# R Ffam; PF00048; IL8; 1.

# R SMART; SM00199; SCY; 1.

# R SMART; SM00199; SCY; 1.

# R SMALL CYTOKINES CC; 1.

# R SMALL INDUCIBLE CYTOKINE A4.

# PROSITE; PS00472; SMALL INDUCIBLE CYTOKINE A4.

# PROSITE; PS0472; SMALL INDUCIBLE CYTOKINE A4.

# PROSITE: PS0472; SMALL INDUCIBLE CYTOKINE A1.

# PROS
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O4_RAT

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O1-OCT-1996 (Rel. 34, Created)

O1-OCT-1996 (Rel. 34, Last sequence update)

Small inducible cytokine A4 precursor (CCL4) (Macrophage

protein 1-beta) (MIP-1-beta).

CCL4 OR SCYA4 OR MIP1B.

Rattus norvegicus (Rat).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Mur
h Similarity 45.7%; Score 242.5; DB 1; Similarity 45.7%; Pred. No. 2.7e-20; 42; Conservative 23; Mismatches 26;
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60 PAVVFVTRKNRQVCANPEKKWVREYINSLEMS

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Q8hyn5 macaca mula
Q8kyq3 macaca mula
Q8hyq3 macaca mula
Q8hyq3 macaca mula
Q9bwa6 gallus gall
Q91zl0 sigmodon hi
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Canis.
                                                            Euteleostomi
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Enomoto A., Kano R., Hasegawa A.;
"molecular cloning of canine RANTES gene.";
Submitted (DEC-2002) to the EMBL/GenBank/DDBJ databases
EMBL; AB098562; BAC53725.1; -.
SEQUENCE 91 AA; 10179 MW; 01D79538CB8148E7 CRC64;
                                                                                                                                            databases
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; Fissipedia, Canidae;
                                                            rtebrata;
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                                                                                                                      Crisman M.V., Howard
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Pred. No. 7.1e-36;
3; Mismatches 14;
                                                                                                                                                                                                                                                               11;
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                                                                                             SEQUENCE FROM N.A.
Takafuji V.A., Sharova L.V., Crisman M.V., Howard "Equus caballus RANTES mRNA.";
Submitted (APR-2002) to the EMBL/GenBank/DDBJ date EMBL; AF506970; AAM34212.1;
InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_ILB.
InterPro; IPR001811; Chemokine_ILB.
Ffam; PF00048; ILB; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL CYTOKINES CC; 1.
PROSITE; PS00472; SMALL CYTOKINES CC; 1.
                                                                                                                                                                                                               NES CC; 1.
AEB253E8CD4ED7FD
                                                                                                                                                                                                                                                 Score 406; DB 6;
Pred. No. 4.2e-40,
4; Mismatches 1:
01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence upd
01-MAR-2003 (TrEMBLrel. 23, Last annotation u
Small inducible cytokine A5 RANTES.
Equus caballus (Horse).
Eukaryota; Metazoa; Chordata; Craniata; Verte
Mammalia; Eutheria; Perissodactyla; Equidae;
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Last sequence up
Last annotation
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Eukaryota; Metazoa; Chordata; C
Mammalia; Eutheria; Carnivora;
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75.8%;
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Similarity 83.5%;
76; Conservative
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01-MAR-2003 (TrEMBLrel. 2
01-MAR-2003 (TrEMBLrel. 2
01-MAR-2003 (TrEMBLrel. 2
RANTES protein.
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69; Conser
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Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
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Felis.
                                                                                                                                                                                                                Kimura T., Kano R., Hasegawa A.;

Kimura T., Kano R., Hasegawa A.;

Kimura T., Kano R., Hasegawa A.;

"molecular cloning of feline RANTES gene.";

Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases.

EMBL; AB083479; BAB88940.1; -.

InterPro; IPR000827; CC_chemkine_sml.

InterPro; IPR001811; Chemokine_IL8.

Pfam; PF00048; IL8; 1.

SMART; SM00199; SCY; 1.

PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.

SEQUENCE 92 AA; 10167 MW; 2E6F087140BA3CE8 CRC64;
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F MW; 744A64BB229194EF CRC64;
                                                                                                Felis silvestris catus (Cat).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata,
Mammalia, Eutheria, Carnivora, Fissipedia, Felidae,
NCBI_TaxID=9685,
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Last annotation update)
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annotation update)
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MEDLINE=21655115; PubMed=11797102;
Hughes S., Haynes A., O'Regan M., Bumstead N.;
Hughes S., Haynes A., O'Regan M., Bumstead N.;
Identification, mapping, and phylogenetic analys
chicken CC chemokines.";
Immunogenetics 53:674-683(2001).
EMBL; AY037859; AAK84432.1; -.
InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_IL8.
Pfam; PF00048; IL8; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL CYTOKINES CC; 1.
SEQUENCE 91 AA; 10154 MW; 744A64BB229194EF CR
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Pred. No. 7.9e-26;
5; Mismatches 22;
update)
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Pred. No. 5.5e-35;
0; Mismatches 14;
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Last
     (TrEMBLrel. 21, (TrEMBLrel. 22,
  01-JUN-2002 (TrE
01-OCT-2002 (TrE
RANTES protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity 47.8
4: Conservative
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                                PRELIMINARY;
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RESULT
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                                                                                                                                                                                                                         SEQUENCE FROM N.A.
Hofmann-Lehmann R., Williams A.L., Swenerton R.K., Li P.-L.,
Rasmussen R.A., Chenine A.-L., McClure H.M., Ruprecht R.M.;
"Quantification of Simian Cytokine and Beta-Chemokine mRNAs using
Real-Time Reverse-Transcriptase Polymerase Chain Reaction: Variations
in Expression during Chronic Primate Lentivirus Infection.";
Submitted (DEC-2001) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 MKVSAARLAVILIATALCAPASASPYSSDT-TPCCFAYIARPLPRAHIKEYFYTSGKCSN
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08-5002 (TrEMBLrel. 21, Created)
01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-JUN-2002 (TrEMBLrel. 22, Last annotation update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
Macrophage inflammatory protein-1 alpha.
Bos taurus (Bovine).
Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Providae; Bovinae; Bos.
                                                                                                                                         eostomi;
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Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
Cercopithecinae; Macaca.
NCBI_TaxID=9544;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 50;
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"Role of chemokines in RSV infection.";
Submitted (JAN-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AY077840; AAL78060.1; -.
InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_IL8.
Pfam; PF00048; IL8; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.
SEQUENCE 93 AA; 10118 MW; 1266BFBFCEE5E8E9 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                     F58463F73E7E900B CRC64;
                                                                    Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                   59.2%; Score 283; DB 6; I 100.0%; Pred. No. 6.3e-26; iive 0; Mismatches 0;
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(TrEMBLrel. 23, Last seq
(TrEMBLrel. 23, Last ann
                                                                 01-MAR-2003 (TrEMBLrel. 23, Las 01-MAR-2003 (TrEMBLrel. 23, Las RANTES (Fragment).
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5789 MW;
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50; Conservative
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                   PRELIMINARY;
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9
                Q8HYN5
Q8HYN5;
01-MAR-2003
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SEQUENCE

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1 MKVSAARLAVILIATALCAPASASPYSSDT-TPCCFAYIARPLPRAHIKEYFYTSGKCSN
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
Chemokine CCL4/MIP-1BETA,
Macaca mulatta (Rhesus macaque).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
NCBI_TaxID=9544;
                                                                                                                                              Euteleostomi;
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(1)
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Basu S., Schaefer T.M., Ghosh M., Fuller C.L., Reinhart T.A.;
Basu S., Schaefer T.M., Ghosh M., Fuller C.L., Reinhart T.A.;
"Comprehensive cloning and sequencing reveals evolutionary
conservation among all groups of rhesus macaque chemokines.";
Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AF449266; AAN76070.1; -.
SEOUENCE 92 AA; 10119 MW; 021CAA371143D12A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
Basu S., Schaefer T.M., Ghosh M., Fuller C.L., Reinhart T.A.;
Basu S., Schaefer T.M., Ghosh M., Fuller C.L., Reinhart T.A.;
"Comprehensive cloning and sequencing reveals evolutionary
conservation among all groups of rhesus macaque chemokines.";
Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AF449267; AAN76071.1; -.
SEOUENCE 92 AA; 10103 MW; 80B2C2071565F809 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 92;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              th Score 252; DB 6; Length 92; Similarity 50.0%; Pred. No. 5.2e-22; 46; Conservative 24; Mismatches 20; Indels
                                                                                                            Macaca mulatta (Rhesus macaque).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele:
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
Cercopithecinae; Macaca.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
(TrEMBLrel. 23, Created)
(TrEMBLrel. 23, Last sequence update)
(TrEMBLrel. 23, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 251.5; DB Pred. No. 6e-22; 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    60 PGVIFLTKRGRQVCADPSKEWVQKYVSDLELS 91
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Werling D.

[1] SEQUENCE

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Q9PWA6

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MKVSAARLAVILIATALCAPASASPYSSD-TTPCCFAYIARPLPRAHIKEYFYTSGKCSN 59
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_ILB.
Pfam; PF00048; ILB; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL CYTOKINES_CC; 1.
SEQUENCE 92 AA; 10334 MW; CF9AAB3D94DCAF79 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
Macrophage inflammatory protein-1b2.
                                                                                                                                                                                         Q91Z65;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
Macrophage inflammatory protein-1 alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 232; DB 11;
Pred. No. 1.2e-19;
B; Mismatches 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         91
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                                                               PAVVFVTRKNROVCANPEKKWVREYINSLEMS
                                                                                                                                                                              92
                                                                                                                                                                              PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             48.5%;
larity 47.8%;
Conservative 1
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                                                                                                                                                                              PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                           Sigmodon.
NCBI_TaxID=42415;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            44;
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Submitted
EMBL; AY07
InterPro;
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Q91Z65
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                         en).
Chordata; Craniata; Vertebrata; Buteleostomi;
leognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                                                                                                                                                                              mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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; Sigmodontinae;
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                                                                                                                                                                                                                                         Hughes S.M., Bumstead N.;
Hughes S.M., Bumstead N.;
Hughes S.M., Bumstead N.;
"Mapping of the gene encoding the chicken homologue of the chemokine SCYA4.";
Submitted (APR-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AF146730; AAD48772.1; -.
HSSP; P13236; 1HUM.
InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_IL8.
Ffam; PF00048; IL8; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.
SEQUENCE 90 AA; 9986 WW; SOAF9679A26751CB CRC64;
                                                                                                                                                                                                                                                             chicken homologue of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
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01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
Macrophage inflammatory protein 1 beta.
MIP-1BETA.
Sigmodon hispidus (Hispid cotton rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CYTOKINES CC; 1.
MW; A34FDE21E6FA9C2E CRC64
                                                                                 update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.

Blanco J.C., Pletneva L.M., Prince G.A.;

"Sigmodon hispidus cytokines, chemokines and inte Submitted (SEP-2001) to the EMBL/GenBank/DDBJ dat EMBL; AF421392; AAL16933.1; -.

HSSP; Q9Y258; 1G2S.

InterPro; IPR000827; CC_chemkine_sml.

InterPro; IPR001811; Chemokine_IL8.

Pfam; PF00048; IL8; 1.

SMART; SM00199; SCY; 1.

PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.

SEQUENCE 92 AA; 10195 MW; A34FDE21E6FA9C2E CH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        51.8%; Score 247.5; DB 11, larity 46.7%; Pred. No. 1.8e-21; Conservative 21; Mismatches 27;
                                                                    update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    52.0%; Score 248.5; DB 13,
larity 50.0%; Pred. No. 1.3e-21;
Conservative 20; Mismatches 23;
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Ο
                                                                    Last sequence up
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                                                        Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRT;
                                                                                                                               Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; C
Archosauria; Aves; Neognathae;
                                                       m m 0
                                                      (TrEMBLrel. 13 (TrEMBLrel. 13 (TrEMBLrel. 20
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                        PRELIMINARY;
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43; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                  FROM N.A
                                                                                                                                                                                                TaxID=9031;
                                      09PWA6;
01-MAY-2000
01-MAY-2000
01-MAR-2002
Chemokine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    46;
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Query Match Best Local

fatches

SEQUENCE

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Query Match Best Local S Matches 43

Sigmodon.

Q91ZL0 Q91ZL0;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sigmodon hispidus (Hispid cotton rat).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Sigmodontinae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nibbs R.J., Barcellos L.F., Townson J.R.,
"Variation in gene copy number of the human chemokines macrophage inflammatory protein-la/CCL3 and macrophage inflammatory protein-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.

Blanco J.C., Pletneva L.M., Prince G.A.;

"Sigmodon hispidus cytokines, chemokines and interferons.";

Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases.

EMBL; AY059407; AAL26704.1; -.

HSSP; Q9Y258; 1G2S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 92;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Euteleostomi; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Brain;
Ishizuka K., Igata-Yi R., Naruse K., Nakashima H., Ohuchi K.,
Katsuragi S., Kin Y., Ohmoto Y., Nomiyama H., Iio M., Miura R.,
Miyakawa T.;
Submitted (AUG-1995) to the EMBL/GenBank/DDBJ databases.
EMBL; D63785; BAA09855.1; -.
HSSP; P13236; 1HUM.
InterPro; IPR000827; CC_chemkine sml.
InterPro; IPR001811; Chemokine_IE8.
InterPro; IPR001811; Chemokine_IE8.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.
Signal.
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                                                                                                       Length
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InterPro; IPR001811; Chemokine_IL8.
Pfam; PF00048; IL8; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.
SEQUENCE 92 AA; 10166 MW; 4C8D01E926CDE882 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
NCBI_TaxID=9606;
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Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                      Q14745 PRELIMINARY; PRT; 80 AA. Q14745; Q14745; 01-NOV-1996 (TrEMBLrel. 01, Created) 01-NOV-1996 (TrEMBLrel. 01, Last sequence update) 01-MAR-2002 (TrEMBLrel. 20, Last annotation update LD78 alpha beta precursor (Fragment). Homo sapiens (Human).
                                                                                                   Query Match
Best Local Similarity 44.6%; Pred. No. 1.8e-19;
Matches 41; Conservative 22; Mismatches 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 213; DB 4; I
Pred. No. 1.7e-17;
21; Mismatches 22;
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LD78 ALPHA BETA.
                                                                                                                                                                                                                                              60 PAVVFVTRKNRQVCANPEKKWVREYINSLEMS 91
                                                                                                                                                                                                                                                                   61 PAVVFQTKRGKQVCADPSESWVQEYVYDLELN 92
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21;
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Q918E0,
01-OCT-2000 (TrEMBLrel. 15, C
01-OCT-2000 (TrEMBLrel. 15, L
01-DEC-2001 (TrEMBLrel. 19, L
Chemokine K203 precursor.
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nilarity 45.6%;
Conservative 2:
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16
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80
8857 MW;
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Best Local S
Matches 36
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                     Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                                                                                                                                                                                                                              .;
?
                                                                                                                                                                                                                                                                                     Length 89;
                                                                                                                                                                                                                                                                                                              30; Indels
                                    SEQUENCE FROM N.A.

MEDLINE=20170941; PubMed=10704244;

SA Sick C., Schneider K., Staeheli P., Weining K.C.;

"Novel chicken CXC and CC chemokines.";

Cytokine 12:181-186(2000).

E EMBL; Y18692; CAB70956.1; -.

R HSSP; P13236; 1HUM.

R HSSP; P13236; 1LB.

R FRO048; ILB; 1.

Signal.

Signal.

Signal.
                                                                                                                                                                                                                                 POTENTIAL.
CHEMOKINE K203.
6FA2EA7A4950CA75 CRC64;
                                                                                                                                                                                                                                                                                    44.6%; Score 213; DB 13;
larity 42.7%; Pred. No. 1.9e-17;
Conservative 19; Mismatches 30;
                                                                                                                                                                                                                                                                                                                                                                                           61 AVVFVTRKNRQVCANPEKKWVREYINSLE 89
                                                                                                                                                                                                                                                                                                                                                                                                       |::|:|:|:|:|:|
59 AIIFITKKEREVCANPSDPWVQRYLQSVK 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                        completed: December 16, 2003, 15:39:37 te : 31.3333 secs
                                                                                                                                                                                                                                  21
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ALIGNMENTS

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RANTES; T-cell; T-lymphocyte; heparanase; heparin; heparan sulfate; arthritis; restenosis; cancer; wound healing.
              AAR70802 standard; Protein; 91 AA
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93US-0136117.
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N-PSDB; AAQ85372.
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                                                                       RANTES protein.
                                                                                                                Homo sapiens.
                                                                                                                                                                                                                      Hoogwerf AJ,
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detecting
Screening for cpds. with anti-heparanase activity - by inhibition of heparin or heparan sulphate degradation, potentially useful for treating arthritis, restenosis,
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60pp; English Claim 13; Page 51; Purified heparanases, prepared under reducing conditions and activated with transglutaminase, are given in AAR70786-802. Most are prepared by reverse transcription of mRNA from activated human leukocytes, then cloning of the cDNA into pVL1392 baculovirus vector, and expression in Sf9 cells in the presence of reduced glutathione and dithiothreitol.
(Updated on 25-MAR-2003 to correct PN field.)

91 AA; Sequence

ô Gaps .. 0 91 Length Indels Score 478; DB 16; Pred. No. 3.1e-46; 0; Mismatches 0; ·, 100.0%; ilarity 100.0%; Conservative 0; Local Similarity les 91; Conser Match uery Ma lest Loc latches

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AAR97665;

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ed; Human; RANTES; Raised on Activation, normal T cell derived and Secrbone marrow; antagonist; MIP-lalpha; asthma; allergic rhinitis;
atopic dermatitis; atheroma; atherosclerosis; rheumatoid arthritis;
inflammation; chemotaxis.

Homo sapiens.

peptide" claim Location/Qualifiers
1..23
/note= "Signal peptic
24..91
/note= "RANTES, claim Peptide Protein

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WO9617935-A2

95WO-GB02861 07-DEC-1995;

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LTD GROUP GLAXO (GLAX)

Wells AEI, Proudfoot

WPI; 1996-287180/29 N-PSDB; AAT30329.

Polypeptide functioning as antagonist to RANTES or to MIP-1-alphaused in the treatment of e.g. asthma, allergic rhinitis, atheroma/atherosclerosis or rheumatoid arthritis

English 33pp; ., H Example; Fig

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     This sequence represents full length human RANTES (Raised on Activation, normal T cell derived and Secreted). The DNA encoding this sequence was cloned from a human bone marrow lambda gtll cDNA library using the primer sequences given in AAT30330-33. RANTES was used in the design of peptides which act as antagonists to RANTES or to MIP-lalpha (see also AAR97666-68). These peptides may be used in the treatment of asthma, allergic rhinitis, atopic dermatitis, atheroma/atherosclerosis or rheumatoid arthritis by inhibiting or reducing inflammation mediated by RANTES or MIP-lalpha. They are also useful in studying RANTES induced chemotaxis, mobilisation of Ca2+ and receptor binding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C-C chemokine; RANTES; MCP2; chemokine antagonist; inflammatory disease HIV infection; tumour; angiogenesis-related disease; autoimmune disease haematopoiesis-related disease; CD26/DPP IV; immune disease; diagnosis; atherosclerosis; pulmonary disease; skin disorder; therapy.
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Pred. No. 3.1e-46;
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                                                                                                                                                                                                                                                            Wild type; C-C chemokine; monocyte chemotactic protein 2; MCP2; HIV; regulated on activation normal T-cell expressed and secreted; RANTES; truncation; antagonist; medicaments; diagnosis; inflammation; infection; tumour; angiogenesis; hematopoiesis; autoimmune disease; atherosclerosis; pulmonary disease; skin disorder.
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MKVSAARLAVILIATALCAPASASPYSSDTTPCCFAYIARPLPRAHIKEYFYTSGKCSNP
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Pred. No. 3.1e-46;
Mismatches 0;
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                                                                     AVVFVTRKNRQVCANPEKKWVREYINSLEMS
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AIDS; viral; RANTES; vMIP; SDF-1; MPAV;

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The invention relates to a cross-over protein produced by chemical ligation of at least two functional protein modules derived from at least two parent protein molecules. The cross-over proteins can be used in two parent protein molecules. The cross-over proteins and infectious diseases including AIDS, and for indications of hematopoiesis and chemoprotection. They are also useful for treatment of asthma, allergic rhinitis, atopic dermatitis and rheumatoid arthritis. A library comprising a collection of cross-over proteins is useful for screening for cross-over proteins that are receptor ligands. The libraries comprise functionally diverse compounds therefore improving the drug discovery process. The proteins and libraries are exemplified by the preparation of cross-over chemokines comprising various combinations of peptide segments derived from RANTES, SDF-1 (stromal cell derived factor 1), vMIP (viral Macrophage Inflammatory protein) and other such chemokines. The present sequence represents a native amino acid fragment of RANTES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New cross-over proteins for treatment of inflammation and infections e.g. AIDS - prepared by ligation of two functional protein modules derived from two different parent molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chemokine; cross-over protein; pharmaceutical; inflammatory; AIDS; infectious disease; hematopoiesis; chemoprotection; asthma; RANTES; allergic rhinitis; atopic dermatitis; rheumatoid arthritis; SDF-1; stromal cell derived factor 1; Macrophage Inflammatory protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 478; DB 20;
Pred. No. 3.1e-46;
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91; Conservative C
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                            AAY06727 standard; protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4; Page 41;
                                                                                                                    Amino acid sequence
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                                                 Macrophage recruitment; chemokine derivative; MCP-1; osteoporosis; monocyte chemoattractant protein-1; inflammation; atherosclerosis; HIV; AIDS; stroke; psoriasis; autoimmune disease; hypertension; endotoxaemia; basophil-mediated disease; myocardial infarction; acute ischaemia; rheumatoid arthritis; contraception.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention concerns the identification of a number of chemokines which can be used to produce derivatives, agonists and antagonists which are then useful in disease treatment. The chemokines include sequences AAB15785-B15794, AAB15803-B15813 and AAB15831-B15848. These chemokine derivatives can be used to treat diseases such as autoimmune diseases, atherosclerosis, osteoporosis, HIV infection and AIDS, psoriasis, inflammatory diseases, hypertension, basophil-mediated diseases, endotoxaemia, myocardial infarction, acute ischaemia and rheumatoid arthritis, and can be used to prevent strokes and as contraceptives. The coding sequences for the chemokines can be used in gene therapy for the same diseases, as well as in the production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                 tumour
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3.1e-46;
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                         SEQ ID NO:
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                          Human chemokine RANTES
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                                                                                                                                           Homo sapiens.
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17-MAR-1999;
01-DEC-1999;
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17-JAN-2001
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Human, vaccination, gene therapy, nutritional supplement, stem cell proliferation, haematopoiesis, nerve tissue regeneration, immune stimmlation, anti-inflammatory, leukaemia
secreted protein
human
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Homo sapiens

WO200179449-A2

25-0CT-2001

2001WO-US08656 16-APR-2001; 18-APR-2000; 2000US-0552929 26-JAN-2001; 2001US-0770160

(HYSE-) HYSEQ INC

Drmanac RT Liu C, Tang YT,

.ч О in genet of human polypeptides, useful Nucleic acids encoding a range o vaccination, testing and therapy

765pp; English. 206; Claim 20; Page The invention relates to novel human secreted polypeptides. The polypeptides and antibodies to the polypeptides are useful for determining the presence of or predisposition to a disease associated with altered levels of polypeptide. The polypeptides are also useful for identifying agents (agonists and antagonists) that bind to them. Cells expressing the proteins are useful for identifying a therapeutic agent for use in treatment of a pathology related to aberrant expression or physiological interactions of the polypeptides and cells genetically the nucleic acids encoding the polypeptides and cells genetically engineered to express them are also useful for producing the proteins. They may be used to therapy, and can be used as nutritional supplements. They may be used in therapy, and can be used as nutritional supplements. They may be used in therapy, and can be used as nutritional supplements, and in bone, cartilage, tendon and/or nerve tissue growth or regeneration; immune suppression and/or stimulation; as anti-inflammatory agents; and in treatment of leukaemias. AAU29510-AAU33304 represent the amino acid sequences of novel human secreted proteins of the invention. for t 0 and

445 AA; Sequence

ö Gaps . 0 Length Indels 100.0%; Score 478; DB 22; larity 100.0%; Pred. No. 1.9e-45; Conservative 0; Mismatches 0; Local Similarity nes 91; Conser Query Match Sest Local S Matches 91

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ULT 9 1240

standard; Protein; AAM24003

AAM24003;

(first entry) 12-OCT-2001

EST encoded protein SEQ ID NO: 1528

macaque; horse; Human; sheep; pig; cow; fruit fly; yeast; hamster; macaque; hotomato; monkey; dog; sea urchin; expressed sequence tag; EST;

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coding sequences of novel
                                                                                                                                                                                                                                                                                                                                                                 The present invention provides the protein and coding sequences of novel proteins from a variety of organisms, including human, dog, cat, horse, cow, pig, hamster, monkey, macaque, yeast, bacteria, fruit fly, sea urchin and tomato. These were derived from expressed sequence tags (ESTs from the organism of interest. They can be used in diagnostics, forensics, gene mapping, identification of mutations, to assess biodiversity and for nutritional purposes. The present sequence is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 MKVSAARLAVILIATALCAPASASPYSSDTTPCCFAYIARPLPRAHIKEYFYTSGKCSNP
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diagnostics; forensic test; gene mapping; genetic disorder biodiversity; gene therapy; nutrition.
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Pred. No. 1.5e-45;
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24..91
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, Zhang J, F
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; 2000US-0617746.
; 2000US-0631451.
; 2000US-0663870.
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98.9%;
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                                                                                                                2001WO-US02687
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        the invention.
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N-PSDB; AAH98662.
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', Drmanac RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                WO200154477-A2.
                                                                                                                                       25-JAN-2000;
17-JUL-2000;
03-AUG-2000;
15-SEP-2000;
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WO200255019-A2
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18-JUL-2002

2001WO-US45244 23-OCT-2001; 24-OCT-2000; 2000US-243264P

(UNIW) UNIV WASHINGTON

Holtzman MJ;

WPI; 2002-566709/60

and treatment children <u>.</u> RANTES chemokine or an expression system in its of paramyxovirus infection, especially in of paramyxovirus diagnosis of

English. -; 19pp; Claim 5; Page

only The invention relates to the treatment of paramyxovirus infection that involves administration of an active ingredient such as RANTES chemokine (I) or an expression system (II). (I) and (II) are used in the treatment and diagnosis of paramyxoviral infection, especially in children. RANTES chemokine has a significant effect on respiratory infections caused by paramyxovirus. RANTES acts downstream of viral entry and signals through specific CCRI and/or CCR5 chemokine receptors to interrupt the death pathway of macrophages which have been infected by virus. RANTES not only inhibits apoptosis of infected macrophage but also clears the macrophage of infection. The present sequence represents a human RANTES protein. Note: the present sequence is not provided in the specification. It has been obtained from the NCBI database under the accession no. NP_002976.

91 AA; Sequence

Gaps .. 0 Length 91; Indels .. H Score 472; DB 23; Pred. No. 1.5e-45;); Mismatches 1 .; 0 Similarity 98.9%; Conservative 90 Query Match Best Local Matches

- 09 9
- 91

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SULT 11 321079

AAO21079 standard; Protein; 91

(first entry) 19-JUL-2002

Wild type RANTES protein.

human; RANTES; neuroprotective; antiallergic; antiinflammatory; anti-HIV; lchemokine mutant; cationic site; multiple sclerosis; HIV infection; inflammatory disease; demyelinating disease; allergic; wild type.

Homo sapiens

WO200228419-A2

11-APR-2002

2001WO-EP11428 03-OCT-2001; 2000EP-0121665 04-OCT-2000; HOLDING SYSTEMS RES APPLIED) ARS (ISTF

Kosco-Vilbois Wells TNC, Proudfoot A,

2002-340073/37. WPI; A mutant of the human CC chemokine RANTES with two mutations in the cationic site of the 40's loop is used for treatment of multiple sclerosis and/or demyelinating diseases -

Examples; Page 36; 46pp; English.

chemokine mutant), comprising the amino sequence of 91 amino acids as given in the specification. The CC chemokine mutant RANTES, with two mutations in the cationic site is useful for the preparation of a pharmaceutical composition used in treating multiple sclerosis or other demyelinating diseases. The mutant with single mutations at cationic sites is used for the treatment of HIV infection and/or other allergic or inflammatory diseases. This sequence represents the wild-type human RANTES protein of the invention. The invention relates to a truncated and mutated human RANTES

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91 AA; Sequence

.. 0 Gaps ·. Length 91; Indels .. H DB 23; Score 472; DB 23; Pred. No. 1.5e-45; 0; Mismatches 1 ô 98.78; 98.98; Similarity 98.8 0; Conservative 90; Query Match Local

91 AVVFVTRKNROVCANPEKKWVREYINSLEMS AVVFVTRKNRQVCANPEKKWVREYINSLEMS

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RESULT AAM52

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AAM52443 standard; Protein; 91 AA

(first entry) 03-JUL-2002 HIV_Nef1 fusion protein #10

Selenoprotein; HIV; Ebola virus; cancer; immune system disorder

Homo sapiens.

US6303295-B1

16-OCT-2001.

96US-0679493. 12-JUL-1996;

95US-001203P. 95US-003112P. 14-JUL-1995; 01-SEP-1995; CS; Ramanathan Nadimpalli RG, Taylor EW,

(UYGE-) UNIV GEORGIA RES FOUND INC

WPI; 2002-024734/03.

human system e G and immune New selenoprotein for use in detecting certain viruses, immunodeficiency virus (HIV) or Ebola, cancer and immun

Disclosure; Columns 73-76; 140pp; English.

The present invention relates to selenoproteins encoded in the genome a virus, where the coding sequence of the selenoprotein is genetically

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engineered for expression in a nucleic acid construct. The invention als discloses a method for identifying selenoprotein coding sequences, for detecting certain viruses (e.g. HIV or Ebola), cancer and immune system disorders. The present sequence was used to illustrate the invention.
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chemokine mutant; cationic site; multiple sclerosis; HIV infection inflammatory disease; demyelinating disease; allergic; mutein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           site of the 40's loop is used for treatment of multi-
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ed. No. 1.5e-45;
Mismatches 1;
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Pred. No. 6.9e-45;
; Mismatches 2.
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Pred. No. :
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Similarity 97.8%;
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Synthetic.
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Pred. No. 6.9e-45;
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RANTES; neuroprotective; antiallergic; antiinflammatory; anti-HIV; } chemokine mutant; cationic site; multiple sclerosis; HIV infection; inflammatory disease; demyelinating disease; allergic; mutein.
                                                                                                           "Wild-type Arg substituted by Ala"
                                                                                              Location/Qualifiers 70
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human;

A mutant of the human CC chemokine RANTES with two mutations in the cationic site of the 40's loop is used for treatment of multiple sclerosis and/or demyelinating diseases -. ₹ Σ Kosco-Vilbois HOLDING Examples; Page 40; 46pp; English. (ISTF) ARS APPLIED RES SYSTEMS TNC, Wells WPI; 2002-340073/37 4 Proudfoot

2001WO-EP11428

03-OCT-2001;

11-APR-2002

WO200228419-A2

2000EP-0121665

04-OCT-2000;

The invention relates to a truncated and mutated human RANTES (a CC chemokine mutant), comprising the amino sequence of 91 amino acids as given in the specification. The CC chemokine mutant RANTES, with two mutations in the cationic site is useful for the preparation of a pharmaceutical composition used in treating multiple sclerosis or other demyelinating diseases. The mutant with single mutations at cationic sites is used for the treatment of HIV infection and/or other allergic or inflammatory diseases. This sequence represents a mutant human RANTES protein of the invention.

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Pred. No. 6.9e-45;
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Sequence 9, Application US/09144838A
Patent No. US20020051996A1
GENERAL INFORMATION:
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2: /cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

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US-08-927-939-21
; Sequence 21, Application US/08927939
; Publication No. US20010006640A1
; GENERAL INFORMATION:
; APPLICANT: Grainger, David J.
; APPLICANT: Tatalick, Lauen Marie
; TITLE OF INVENTION: Compounds and methods to in;
TITLE OF INVENTION: augment an inflammatory res;
FILE REFERENCE: 295.022US1
; CURRENT APPLICATION NUMBER: US/08/927,939
; CURRENT FILING DATE: 1997-09-11
; NUMBER OF SEQ ID NOS: 83
; SOFTWARE: FastSEQ for Windows Version 3.0
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Sequence 29, Application US/09834795A
Patent No. US20020076710A1
GENERAL INFORMATION:
APPLICANT: Lawrence, Papsidero
APPLICANT: Lyn, Dyster
APPLICANT: Lyn, Dyster
APPLICANT: Jana, Frustaci
TITLE OF INVENTION: Detection and Treatment of Breast Cancer
FILE REFRENCE: 3380/11127-US3
CURRENT APPLICATION NUMBER: US/09/834,795A
CURRENT FILING DATE: 1998-09-03
PRIOR FILING DATE: 1998-09-03
PRIOR FILING DATE: 1998-01-20
PRIOR FILING DATE: 1998-01-20
PRIOR FILING DATE: 1998-07-09
NUMBER OF SEQ ID NOS: 35
SOFTWARE: Patentin version 3.0
APPLICANT: Siani, Michael A.
APPLICANT: Wilken, Jill
APPLICANT: Wilken, Jill
APPLICANT: Simon, Reyna
APPLICANT: Simon, Reyna
APPLICANT: Simon, Reyna
APPLICANT: Stephen B.H.
TITLE OF INVENTION: Modular Protein Libraries and Methods of Pre
FILE REFERENCE: GRFN-020/01US
CURRENT APPLICATION NUMBER: US/09/144,838A
CURRENT FILING DATE: 1998-08-31
EARLIER APPLICATION NUMBER: US 60/057,620
EARLIER FILING DATE: 1997-09-04
NUMBER OF SEQ ID NOS: 54
SOFTWARE: Patentin Ver. 2.1
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Best Local
Matches 9
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; Sequence 8, Application US/09920137A
; Publication No. US20030049725A1
; GENERAL INFORMATION:
; APPLICANT: Coleman, Roger
; APPLICANT: Bandman, Olga
; APPLICANT: Wilde, Craig G.
; TITLE OF INVENTION: NEW CHEMOKINES EXPRESSED IN PANCREAS
; NUMBER OF SEQUENCES: 11
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Incyte Pharmaceuticals, Inc.
; STREET: 3174 Porter Drive
; CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                        Length 91;
                                                                                                   and Treatment of Breast Cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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Pred. No. 6.4e-47;
Mismatches 0;
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APPLICANT: Lyn, Dyster
APPLICANT: Lyn, Dyster
APPLICANT: Jana, Frustaci
TITLE OF INVENTION: Detection and Treatment of FILE REFERENCE: 3380/11127-US4
CURRENT APPLICATION NUMBER: US/09/834,794A
CURRENT FILING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: 09/146,580
PRIOR FILING DATE: 1998-01-20
PRIOR FILING DATE: 1998-01-20
PRIOR FILING DATE: 1998-01-20
PRIOR FILING DATE: 1998-07-09
NUMBER OF SEQ ID NOS: 35
SOFTWARE: PatentIn version 3.0
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Application US/09834794A
No. US20030026777A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/920,
FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Sc
100.0%; Pr
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                        : Homo sapiens
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                                 INFORMATION:
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Sequence 29, App
Publication No.
GENERAL INFORMAT
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                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM:
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Best Local
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Sequence 1, Application US/09537858

Publication No. US20030119148A1

SENERAL INFORMATION:
APPLICANT: PROOST, PAUL
APPLICANT: STRUYF, SOFIE
APPLICANT: VAN DAMME, JO
TITLE OF INVENTION: AMINO-TERMINALLY TRUNCATED RANTES AS CHE
TITLE OF INVENTION: ANTAGONISTS
FILE REFERENCE: 2024/49674
CURRENT APPLICATION NUMBER: US/09/537,858
CURRENT FILING DATE: 2001-02-26
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
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Sequence 8, Application US/10137655
Publication No. US20030138917A1
GENERAL INFORMATION:
APPLICANT: Coleman, Roger
APPLICANT: Bandman, Olga
APPLICANT: Wilde, Craig G.
TITLE OF INVENTION: NEW CHEMOKINES EXPRESSED IN NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         h 100.0%; Score 478; DB 11; Similarity 100.0%; Pred. No. 6.4e-47; 91; Conservative 0; Mismatches 0;
                                                                                                                                     Score 478; DB 11;
Pred. No. 6.4e-47;
); Mismatches 0;
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                                                                                                                                      100.0%;
larity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                               TOPOLOGY: linear
MOLECULE TYPE: peptide
IMMEDIATE SOURCE:
LIBRARY: GenBank
CLONE: RANTES
                      single
TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                       Similarity
91; Conser
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-10-137-655-8
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Best Local S
Matches 91
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Sest Local S
Watches 91
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Wilde, Craig C.
Seilhamer, Jeffrey J.
TITLE OF INVENTION: CHEMOKINE EXPRESSED IN FETAL SPLEEN,
ITS PRODUCTION AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 478; DB 12;
Pred. No. 6.4e-47;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/158,366
FILING DATE: 29-May-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/467,123B
FILING DATE: 06-JUN-1995
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                                                                          OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/137,655
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0027 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEPAX: 415-852-0195
INFORMATION FOR SEQ ID NO: 8:
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; Sequence 5, Application US/10158366
; Publication No. US20020142398A1
; GENERAL INFORMATION:
; APPLICANT: Coleman, Roger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Sc
100.0%; Pr
                                             IBM Compatible
                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                 peptide
                               COMPUTER READABLE FORM:
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COUNTRY: US
ZIP: 94304
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RANTES
                                          MEDIUM TYPE: Dist
COMPUTER: IBM CON
OPERATING SYSTEM:
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LIBRARY: GenBa
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COUNTRY:
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SEQUENCE DESCRIPTION: SEQ ID NO: 8:
MOLECULE TYPE: peptide
IMMEDIATE SOURCE:
LIBRARY: GenBank
CLONE: RANTES
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US-10-293-705-12
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                                                                                US-10-057-275-8
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LENGTH: 91
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Best Local 9
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                      Length 91;
 APPLICATION NUMBER: US 08/375,346
FILING DATE: 19-JAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0026-1 DIV
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-555-0555
TELEPHONE: 415-845-4166
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
                                                                                                                                                                                                                                                                                                    Score 478; DB 14;
Pred. No. 6.4e-47;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 94304

ZIP: 94304

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/057,275
FILING DATE: 25-Jan-2002
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ALIUANEI/ AGENTA .... Barbara J.
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0027 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/390,740A
FILING DATE: February 17, 1995
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                         ID NO:
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Publication No. US20020155545A1
GENERAL INFORMATION:
APPLICANT: Coleman, Roger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
                                                                                                                                                                                                                                                                                                   100.0%; Silarity 100.0%; EConservative 0;
                                                                                                                                                                                                                                                          SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                   TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: 5
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                                                                                                                                                                                                                                                                                                                        Similarity
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 Length 91;
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APPLICANT: Wilde, Craig G.
APPLICANT: Hawkins, Phillip R.
APPLICANT: Bandman, Olga
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: DNA ENCODING LIVER EXPRESSED CHEMOKINE
FILE REFERENCE: PF-0024-3 CON
CURRENT APPLICATION NUMBER: US/10/293,705
CURRENT FILING DATE: 2002-11-12
PRIOR FILING DATE: 1998-12-09
PRIOR FILING DATE: 1998-12-09
PRIOR FILING DATE: 1994-11-29
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PERL Program
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Siani, Michael A.
APPLICANT: Wilken, Jill
APPLICANT: Simon, Reyna
APPLICANT: Kent, Stephen B.H.
TITLE OF INVENTION: Modular Protein Libraries and Methods of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature
OTHER INFORMATION: GenBank ID No. US20030083468A1 g134510
                                        Indels
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larity 100.0%; Pred. No. 6.4e-47;
Conservative 0; Mismatches 0;
100.0%; Score 478; DB 14;
100.0%; Pred. No. 6.4e-47;
tive 0; Mismatches 0;
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Publication No. US20030083468A1
GENERAL INFORMATION:
                                        Conservative
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ORGANISM: Homo sapiens
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Sequence 42, Application US/09144838A
Sequence 42, Application US/09144838A
Batent No. US20020051996A1
GENERAL INFORMATION:
APPLICANT: Siani, Michael A.
APPLICANT: Siani, Michael A.
APPLICANT: Simon, Reyna
APPLICANT: Kent, Stephen B.H.
TITLE OF INVENTION: Modular Protein Libraries and Methods of Preparation
TITLE OF INVENTION: Modular Protein Libraries and Methods of Preparation
CURRENT APPLICATION NUMBER: US/09/144,838A
CURRENT FILING DATE: 1998-08-31
EARLIER FILING DATE: 1997-09-04
NUMBER OF SEQ ID NOS: 54
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 42
LENGTH: 68
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Pred. No. 3.2e-35;
0; Mismatches 0;
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Pred. No. 3.2e-35;
0; Mismatches 0;
FILE REFERENCE: GRFN-020/01US
CURRENT APPLICATION NUMBER: US/09/144,838A
CURRENT FILING DATE: 1998-08-31
EARLIER APPLICATION NUMBER: US 60/057,620
EARLIER FILING DATE: 1997-09-04
NUMBER OF SEQ ID NOS: 54
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 10
LENGTH: 68
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Similarity 100.0%; E
68; Conservative 0;
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Best Local Similarity 100.0%; E
Matches 68; Conservative 0;
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Patent No. US20020081623A1
GENERAL INFORMATION:
APPLICANT: WILLIAMS, TIMOTHY
APPLICANT: JOSE, PETER J.
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US-09-792-793A-29

US-09-792-793A-29

Sequence 29, Application US/09792793A

Patent No. US20020168370A1

GENERAL INFORMATION:

APPLICANT: McDonald, John R.

APPLICANT: Coggins, Philip

TITLE OF INVENTION: OTHER INFLAMMATORY CONDITIONS AND DISORDERS

FILE REFERENCE: 25020-601D

CURRENT APPLICATION NUMBER: US/09/792,793A

CURRENT FILING DATE: 2001-02-22

NUMBER OF SEQ ID NOS: 93

SOFTWARE: Patentin Ver. 2.0
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US-09-792-793A-29
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; Pred. No. 3.2e-35;
0; Mismatches 0;
APPLICANT: GRIFFITHS-JOHNSON, DAVID A.
APPLICANT: HSUAN, JOHN J.
TITLE OF INVENTION: CHEMOTACTIC CYTOKINE
FILE REFERENCE: 550-33
CURRENT APPLICATION NUMBER: US/09/195,457
CURRENT FILING DATE: 1998-11-18
PRIOR APPLICATION NUMBER: 08/470,323
PRIOR FILING DATE: 1995-06-06
PRIOR FILING DATE: 1994-09-14
PRIOR APPLICATION NUMBER: GB 9318984.3
PRIOR FILING DATE: 1993-09-14
PRIOR FILING DATE: 1993-09-14
PRIOR FILING DATE: 1994-04-29
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US-10-375-209A-29
; Sequence 29, Application US/10375209A
; Publication No. US20030215421A1
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ORGANISM: homo sapien
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GENERAL INFORMATION:
APPLICANT: McDonald, John R.
APPLICANT: Coggins, Philip
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING SECONDARY TISSUE DAMAGE AND
TITLE OF INVENTION: OTHER INFLAMMATORY CONDITIONS AND DISORDERS
FILE REFERENCE: 25020-601E
CURRENT APPLICATION NUMBER: US/10/375,209A
CURRENT FILING DATE: 2003-02-24
NUMBER OF SEQ ID NOS: 93
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 29
LENGTH: 68
TYPE: PRT
ORGANISM: homo sapien
FEATURE:
OTHER INFORMATION: Human Chemokine Polypeptide: RANTES
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; Sequence 36, Application PC/TUS9408207A
; GENERAL INFORMATION:
; APPLICANT: The Upjohn Company
; TITLE OF INVENTION: USE OF HEPARANASE TO IDENTIFY AND
; TITLE OF INVENTION: ISOLATE ANTI-HEPARANASE COMPOUNDS
; NUMBER OF SEQUENCES: 43
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: The Upjohn Company, Intellectual Property La
STREET: 301 Henrietta
; CITY: Kalamazoo
; STATE: MI
; COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ט
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Version
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vers:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/08207A
FILING DATE:
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DZ,

ult No.

Dec

amino acids

LENGTH:

peptide

MOLECULE TYPE:

JT-US94-08207-36

single

TYPE: amino acid STRANDEDNESS: sir TOPOLOGY: linear

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APPLICANT: Hoogewerf, Arlene J.
APPLICANT: Ledbetter, Steven R.
TITLE OF INVENTION: USE OF HEPARANASE TO IDENTIFY AND
TITLE OF INVENTION: ISOLATE ANTI-HEPARANASE COMPOUNDS
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: The Upjohn Company, Intellectual Property Law
STREET: 301 Henrietta
CITY: Kalamazoo
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            Length 91;
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                                        Indels
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 th 100.0%; Score 478; DB 5; Similarity 100.0%; Pred. No. 2.9e-46; 91; Conservative 0; Mismatches 0;
            Score 478; DB 1;
Pred. No. 2.9e-46;
; Mismatches 0;
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GENERAL INFORMATION:
APPLICANT: Coleman, Roger
                                                                                                                                                                                                                          US-08-136-117-36
; Sequence 36, Application US/08136117
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Jameson, William G.
REGISTRATION NUMBER: 27,199
REFERENCE/DOCKET NUMBER: 4731
TELECOMMUNICATION INFORMATION:
TELEPHONE: 616/385-7561
TELEFAX: 616/385-6897
100.0%; Sc.
100.0%; Pr
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 91 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                           Conservative
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FILING DATE:
CLASSIFICATION: 514
                               Similarity
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US-08-390-740B-8
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-08-467-123-5
Sequence 5, Application US/08467123
GENERAL INFORMATION:
APPLICANT: Coleman, Roger
APPLICANT: Wilde, Craig G.
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: A NEW CHEMOKINE EXPRESSED IN FETAL SPLEEN, TITLE OF INVENTION: ITS PRODUCTION AND USES
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
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APPLICANT: Bandman, Olga
APPLICANT: Wilde, Craig G.
TITLE OF INVENTION: NEW CHEMOKINES EXPRESSED IN PANCREAS
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 478; DB 7;
Pred. No. 2.9e-46;
; Mismatches 0;
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                                                                                                                                                                                                           COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/390,740B
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0027 US
TELEPHONE: 415-855-0195
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Diskette
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version
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100.0%;
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         single
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STRANDEDNESS: si
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CLONE: RANTES
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Best Local Similarity
Matches 91; Conser
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GENERAL INFORMATION:
APPLICANT: Grainger, David J.
APPLICANT: Tatalick, Lauen Marie
TITLE OF INVENTION: Compounds and methods to inhibit or;
TITLE OF INVENTION: augment an inflammatory response;
FILE REFERENCE: 295.022US1;
CURRENT APPLICATION NUMBER: US/08/927,939;
CURRENT FILING DATE: 1997-09-11
NUMBER OF SEQ ID NOS: 83;
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     methods to inhibit or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 478; DB 8;
larity 100.0%; Pred. No. 2.9e-46;
Conservative 0; Mismatches 0;
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Pred. No. 2.9e-46;
; Mismatches 0;
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      APPLICATION NUMBER: US/08/467,123
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/375,346
FILING DATE: 19-JAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: LUTHER, BARBARA J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0026 US
TELEPHONE: (415) 855-0555
TELEFAX: (415) 855-0572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 AVVFVTRKNRQVCANPEKKWVREYINSLEMS
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100.0%; Pr
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TYPE: amino acids
STRANDEDNESS: single
                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                 internal
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                  linear
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ORIGINAL SOURCE:
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US-08-927-939-21
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LENGTH: 91
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Best Local (
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US-09-023-092-4
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Sequence 4, Application US/09023092
GENERAL INFORMATION:
APPLICANT: Hedrick, Joseph A.
APPLICANT: Wang, Luquan
APPLICANT: Wang, Luquan
APPLICANT: Albert
APPLICANT: Albert
APPLICANT: And Mammalian Chemokines; Related Reagents
TITLE OF INVENTION: And Methods
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: DNAX Research Institute
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Pred. No. 2.9e-46;
Mismatches 0;
                  -08-927-939A-21
Sequence 21, Application US/08927939A
GENERAL INFORMATION:
APPLICANT: Grainger, David J.
APPLICANT: Tatalick, Lauren Marie
TITLE OF INVENTION: Compounds and methods to inl
TITLE OF INVENTION: augment an inflammatory ref
FILE REFERENCE: 1543.001US1
CURRENT APPLICATION NUMBER: US/08/927,939A
CURRENT FILING DATE: 1997-11-09
NUMBER OF SEQ ID NOS: 85
SOFTWARE: Fast SEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/023,092
FILING DATE: 13-FEB-1998
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Ching, Edwin P.
REGISTRATION NUMBER: 34,090
REFERENCE/DOCKET NUMBER: DX0780
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNAX Research Institute
| California Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 94304-1104
ZIP: 94304-1104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TELEFAX: (650)496-1200
RMATION FOR SEQ ID NO: 4
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08-927-939A-21
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Best Local Simi
Matches 91;
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Length 91;
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                          Indels
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SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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Score 478; DB 14;
Pred. No. 2.9e-46;
; Mismatches 0;
                                                                                                                                                                                                 Sequence 8, Application US/09113705
; GENERAL INFORMATION:
; APPLICANT: BANDMAN, OLGA
; APPLICANT: COLEMAN, ROGER
; APPLICANT: STUART, SUSAN G.
; TITLE OF INVENTION: NEW CHEMOKINE EXPRESSED;
; NUMBER OF SEQUENCES: 9
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           91
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APPLICATION NUMBER: US/08/421,144
FILING DATE: 13-APR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US/09/113,705
                                                                                                                                                                                                                                                                                                                                                                              ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
FORMATION FOR SEQ ID NO: 8:
100.0%; S
llarity 100.0%; P
Conservative 0;
                                                                                                                                                                                                                                                                                                                    3174 Porter Drive
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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Rest Local Similarity
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                Similarity
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; MOLECULE TYPE:
US-09-113-705-8
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   Query Match
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TELECOMMUNICATION INFORMATION:
GENERAL INFORMATION:
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ENGTH: 91
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Matches 91
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Pred. No. 2.9e-46;
; Mismatches 0;
                                                                                                                                                                                                                                                                 Score 478; DB 15;
Pred. No. 2.9e-46;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 9, Application US/09144838A
GENERAL INFORMATION:
APPLICANT: Siani, Michael A.
APPLICANT: Wilken, Jill
APPLICANT: Wilken, Jill
APPLICANT: Kent, Stephen B.H.
TITLE OF INVENTION: Modular Protein Libraries and
FILE REFERENCE: GRFN-020/01US
CURRENT APPLICATION NUMBER: US/09/144,838A
CURRENT FILING DATE: 1998-08-31
EARLIER APPLICATION NUMBER: US 60/057,620
EARLIER FILING DATE: 1997-09-04
NUMBER OF SEQ ID NOS: 54
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                      1 MKVSAARLAVILIATALCAPASASPYSSDTTPCCFAY
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Sequence 5, Application US/09113705A
GENERAL INFORMATION:
APPLICANT: Bandman, Olga
APPLICANT: Coleman, Roger
APPLICANT: Stuart, Susan G.
TITLE OF INVENTION: NEW CHEMOKINE EXPRESSED I
FILE REFERENCE: PF-0031-1 DIV
CURRENT APPLICATION NUMBER: US/09/113,705A
CURRENT FILING DATE: 1998-07-10
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                              EXPRESSED
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ce 21, Application US/0915081
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ilarity 100.0%;
Conservative
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                                                                                                                                                                                                    sapiens
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91; Conserv
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91; Conser
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3-09-144-838-9
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                                                                                                                                             SOFTWARE.
SEQ ID NO 5
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Best Local
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APPLICANT: Grainger, David J.
APPLICANT: Grainger, Lauen Marie
APPLICANT: Tatalick, Lauen Marie
APPLICANT: Tatalick, Lauen Marie
TITLE OF INVENTION: Compounds and methods to inhibit or
TITLE OF INVENTION: augment an inflammatory response.
FILE REFERENCE: 295.027US1
CURRENT APPLICATION NUMBER: US/09/150,813
CURRENT FILING DATE: 1998-09-11
BARLIER APPLICATION NUMBER: US 08/927939
EARLIER FILING DATE: 1997-09-11
NUMBER OF SEQ ID NOS: 105
SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Hawkins, Phillip R.
APPLICANT: Bandman, Olga
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: EXPRESSED CHEMOKINES, THEIR
TITLE OF INVENTION: PRODUCTION AND USES
NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                       Score 478; DB 15;
Pred. No. 2.9e-46;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AVVFVTRKNRQVCANPEKKWVREYINSLEMS 91
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals,
STREET: 3174 Porter Drive
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US/09/208,803
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TOMBER: 08/347,492

29-NOV-1994

TOMBER: 08/303,241

07-SEP-1994
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APPLICATION NUMBER: 08/320,011
FILING DATE: 05-OCT-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 13
US-09-208-803-12
; Sequence 12, Application US/09208803
; GENERAL INFORMATION:
; APPLICANT: Wilde, Craig G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08/798,143
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OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/208,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Luther, Barbara J
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF
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PRIOR APPLICATION:
APPLICATION NUMBER: 06
FILING DATE: 10-FEB-15
APPLICATION NUMBER: 06
FILING DATE: 29-NOV-15
APPLICATION NUMBER: 06
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo sapiens -09-150-813-21
                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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APPLICANT: Coleman, Craig C.
APPLICANT: Wilde, Craig C.
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: CHEMOKINE EXPRESSED IN FETAL SPLEEN, TITLE OF INVENTION: ITS PRODUCTION AND USES
NUMBER OF SEQUENCES: 9
                                                                                                                                                                       91;
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pred. No. 2.9e-46;
; Mismatches 0;
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Pred. No. 2.9e-46;
0; Mismatches 0;
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9-360-290-5

quence 5, Application US/09360290

NERAL INFORMATION:
PPLICANT: Bandman, Olga
PPLICANT: Coleman, Roger
PPLICANT: Stuart, Susan G.
ITLE OF INVENTION: NEW CHEMOKINE EXPRESED
ITLE OF INVENTION: NEW CHEMOKINE EXPRESSED
ITLE OF INVENTION: NEW CHEMOKINE EXPRESSED
ITLE REFERENCE: PF-0031-2 DIV
ILE REFERENCE: PF-0031-2 DIV
INRENT FILING DATE: 1999-07-22
IURRENT FILING DATE: 1999-07-22
IURRENT FILING DATE: 1995-04-13
IARLIER FILING DATE: 1995-04-13
IUMBER OF SEQ ID NOS: 8
SOFTWARE: FASTSEQ Version 3.0
                                                                                                                                                                                                                                                                             -09-387-620-5
Sequence 5, Application US/09387620
GENERAL INFORMATION:
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sest Local Similarity 100.0%;
fatches 91; Conservative
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ilarity 100.0%;
Conservative 0
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-360-290-5
                               91 amino acids
STRANDEDNESS: F'TOPOLOGY'
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
ORMATION FOR SEQ ID NO:
EQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
                                                                                                  peptide
                                                                                            TOPOLOGY: linear IOLECULE TYPE: pept: MMEDIATE SOURCE: LIBRARY: GENBANK CLONE: GI 134510
                                                                                                                                                                                              l Similarity
91, Conserv
                                                                                                                                                        -208-803-12
                                                     LENGTH:
TYPE: al
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Pred. No. 2.9e-46;
; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                   PF-0026-1 DIV
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
CITY: Palo Alto
                                                                                                                                              OPERATING SYSTEM: DOS
SOFTWARE: FASESEQ Version 1.5
SOFTWARE: FASESEQ VERSION 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/387,620
FILING DATE:
CLASSIFICATION:
APPLICATION NUMBER: US 08/467,123
APPLICATION NUMBER: US 08/467,123
APPLICATION NUMBER: PF-0026-1 I
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0026-1 I
RELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO: 5:
LENGTH: 01 amino 2012
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TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
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US-10-474-794-277

US-10-474-794-279

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US-60-487-610-2476

US-60-487-610-2330

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ALIGNMENTS

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RESULT 1
US-10-398-457-1
i Sequence 1, Application US/10398457
i Sequence 1, Application US/10398457
i GENERAL INFORMATION:
i APPLICANT: APPLIED RESEARCH SYSTEMS ARS HOLDING N.V.
i TITLE OF INVENTION: CHEMOKINES MUTANTS IN THE TREATMENT OF MULTIPLE SCLEROSIS
i CURRENT APPLICATION NUMBER: US/10/398,457
i CURRENT FILING DATE: 2003-11-13
i NUMBER OF SEQ ID NOS: 37
i SOFTWARE: Patentin version 3.0
i SEQ ID NO 1
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US-60-487-610-2474

Sequence 2474, Application US/60487610

GENERAL INFORMATION:

APPLICANT: CARGILL, Michele

APPLICANT: HUANG, Hongjin

TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH

TITLE OF INVENTION: METHODS OF DETECTION AND USES THEREOF

FILE REFERENCE: CL001469

CURRENT APPLICATION NUMBER: US/60/487,610

CURRENT FILING DATE: 2003-07-17

NUMBER OF SEQ ID NOS: 97101

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 2474

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ORGANISM: Escherichia coli
FEATURE:
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Sequence 1561, Application US/60485450
GENERAL INFORMATION:
APPLICANT: CARGILL, Michele
APPLICANT: CHANG, Sheng-Yung
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: THEREOF
TITLE OF SEQUENCE: CL001470
CURRENT APPLICATION NUMBER: US/60/485,450
CURRENT FILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 47859
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1561
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Sequence 8, Application US/10398457
GENERAL INFORMATION:
APPLICANT: APPLIED RESEARCH SYSTEMS ARS HOLDING N.V.
TITLE OF INVENTION: CHEMOKINES MUTANTS IN THE TREATMENT OF MULTIPLE FILE REFERENCE: WO465
CURRENT APPLICATION NUMBER: US/10/398,457
CURRENT FILING DATE: 2003-11-13
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.0
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US-10-398-457-10; Application US/10398457; Sequence 10, Application US/10398457; GENERAL INFORMATION: GENERAL INFORMATION: APPLICANT: APPLIED RESEARCH SYSTEMS ARS HOLDING N.V. TITLE OF INVENTION: CHEMOKINES MUTANTS IN THE TREATMENT OF MULTIPLE SCLEROSIS; TITLE OF INVENTION: CHEMOKINES MUTANTS IN THE TREATMENT OF MULTIPLE SCLEROSIS
                                                                                                                                                                                                                                 Sequence 9, Application US/10398457
GENERAL INFORMATION:
APPLICANT: APPLIED RESEARCH SYSTEMS ARS HOLDING N.V.
TITLE OF INVENTION: CHEMOKINES MUTANTS IN THE TREATMENT OF MULTIPLE SCLEROSIS
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                                             Gaps
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Pred. No. 3e-45;
0; Mismatches
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larity 97.8%; Pred. No. 3e-45;
Conservative 0; Mismatches
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CURRENT APPLICATION NUMBER: US/10/398,457
CURRENT FILING DATE: 2003-11-13
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                           FILE REFERENCE: WO465
CURRENT APPLICATION NUMBER: US/10/398,457
CURRENT FILING DATE: 2003-11-13
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.0
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llarity 97.8%;
Conservative (
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89; Conserv
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89; Conser
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US-10-398-457-10
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NAME/KEY: SIGNAL
LOCATION: (1)..(23)
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SENERAL INFORMATION:
APPLICANT: APPLIED RESEARCH SYSTEMS ARS HOLDING N.V.
TITLE OF INVENTION: CHEMOKINES MUTANTS IN THE TREATMENT OF MULTIPLE
FILE REFERENCE: WO465
CURRENT APPLICATION NUMBER: US/10/398,457
CURRENT FILING DATE: 2003-11-13
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.0
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Pred. No. 6.7e-44;
); Mismatches 4
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Pred. No. 3e-45;
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Mismatches
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CURRENT FILING DATE: 2003-11-13
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.0
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Best Local Similarity
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89; Conser
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US-10-398-457-13
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NAME/KEY: SIGNAL
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ENGTH: 91
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3-10-398-457-12
Sequence 12, Application US/10398457
GENERAL INFORMATION:
APPLICANT: APPLIED RESEARCH SYSTEMS ARS HOLDING N.V.
TITLE OF INVENTION: CHEMOKINES MUTANTS IN THE TREATMENT OF MULTIPLE SCLEFILE REFERENCE: WO465
CURRENT APPLICATION NUMBER: US/10/398,457
CURRENT FILING DATE: 2003-11-13
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.0
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                                                 SULT 7
3-10-398-457-11
Sequence 11, Application US/10398457
GENERAL INFORMATION:
APPLICANT: APPLIED RESEARCH SYSTEMS ARS HOLDING N.V.
TITLE OF INVENTION: CHEMOKINES MUTANTS IN THE TREATMENT OF MULTIPLE SCLIFILE REFERENCE: WO465
CURRENT APPLICATION NUMBER: US/10/398,457
CURRENT FILING DATE: 2003-11-13
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.0
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Pred. No. 3e-45;
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ilarity 97.8%;
Conservative (
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89; Conser
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NAME/KEY: SIGNAL
LOCATION: (1)..(23)
S-10-398-457-12
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3-10-398-457-11
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NAME/KEY: SIGNAL
LOCATION: (1)..(
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S-10-398-457-3
Sequence 3, Application US/10398457
GENERAL INFORMATION:
APPLICANT: APPLIED RESEARCH SYSTEMS ARS HOLDING N.V.
TITLE OF INVENTION: CHEMOKINES MUTANTS IN THE TREATMENT OF MULTIPLE SCLEROSIS
FILE REFERENCE: WO465
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TITLE OF INVENTION: CHEMOKINES MUTANTS IN THE TREATMENT OF MULTIPLE SCLEROSIS
FILE REFERENCE: WO465
CURRENT APPLICATION NUMBER: US/10/398,457
CURRENT FILING DATE: 2003-11-13
NUMBER OF SEQ ID NOS: 37
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Gaps

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Indels

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COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/622,134
FILING DATE: 18-Jul-2003
FILING DATE: 18-Jul-2003
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Pred. No. 4.5e-35;
                                                                                                                                                                                                                                                        HSUAN, JOHN J.

HITLE OF INVENTION: CHEMOTACTIC CYTOKINE
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: NIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD, BTH FLOOR
CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/291,038

FILING DATE: 14-Apr-1999

FILING DATE: cunknown>
FILING DATE: cunknown>
FILING DATE: cunknown>
FILING DATE: 13-AUG-1996

FILING DATE: 13-AUG-1996

FILING DATE: 14-SEP-1993

FILING DATE: 14-SEP-1993

FILING DATE: 29-APR-1994

FILING DATE: 29-APR-1994
red. No. 4.5e-35;
Mismatches 0;
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sequence Description: SEQ ID NO: 11:
US-10-622-134-11
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NAME: WILSON, MARY J.
REGISTRATION NUMBER: 32,955
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GRIFFITHS-JOHNSON, DAVID A.
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     Pred. No.
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Similarity 100.0%; F
18; Conservative 0;
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TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                     Floppy disk
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ENCE CHARACTERISTICS:
LENGTH: 68 amin
                                                                                                                                                                                         US-10-622-134-11; Application US/10622134; Sequence 11, Application US/10622134; GENERAL INFORMATION:
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COUNTRY: U.S.A.
ZIP: 22201-4714
UTER READABLE FORM:
MEDIUM TYPE: Floppy
         larity 100.0%; P
Conservative 0;
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Best Local
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equence 4, Application PC/TUS0328745
ENERAL INFORMATION:
APPLICANT: CHEMOKINE THERAPEUTICS CORP.
TITLE OF INVENTION: DESIGN OF CHEMOKINE ANALOGS FOR THE TREATMENT OF HUMAN DISEASES
FILE REFERENCE: 59296.00003
FILE REFERENCE: 59296.00003
FILE REFERENCE: 59296.00003
CURRENT APPLICATION NUMBER: PCT/US03/28745
NUMBER OF SEQ ID NOS: 1640
SOFTWARE: Patentin version 3.2
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398-457-7
lence 7, Application US/10398457
lence 7, Application US/10398457
lence 7, Application US/10398457
lence 7, Application:
SRAL INFORMATION:
CLE APPLICANT: APPLIED RESEARCH SYSTEMS ARS HOLDING N.V.

FLE OF INVENTION: CHEMOKINES MUTANTS IN THE TREATMENT OF MULTIPLE SCLEROSIS
LE REFERENCE: W0465
LE REFERENCE: W0465
RRENT APPLICATION NUMBER: US/10/398,457
RRENT FILING DATE: 2003-11-13
RRENT FILING DATE: 2003-11-13
RBER OF SEQ ID NOS: 37
RBER OF SEQ ID NOS: 37
FTWARE: Patentin version 3.0
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Pred. No. 6.7e-44;
); Mismatches 4;
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Similarity 94.6%;
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JULT 15
-10-332-038A-2
Sequence 2, Application US/10332038A
JENERAL INFORMATION:
APPLICANT: Gryphon Therapeutics, Inc.
APPLICANT: Offord, Robin
APPLICANT: Gaertner, Hubert
APPLICANT: Gaertner, Hubert
APPLICANT: Gaertner, Hubert
APPLICANT: Hartley, Oliver
TITLE OF INVENTION: Chemokine Receptor Modulators, Production and Use
FILE REFERENCE: 03504.271
CURRENT APPLICATION NUMBER: US/10/332,038A
CURRENT FILING DATE: 2003-05-06
PRIOR APPLICATION NUMBER: US 60/217,683
PRIOR PLING DATE: 2000-07-12
NUMBER OF SEQ ID NOS: 28
SOFTWARE: PATENTIN Version 3.2
LENGTH: 68
IYPE: PRT
ORGANISM: Homo Sapiens
84 YINSLEMS 91
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61 YINSLEMS 68
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Gaps . 0 68; Indels Length Query Match
Best Local Similarity 100.0%; Pred. No. 4.5e-35;
Matches 68; Conservative 0; Mismatches 0; -10-332-038A-2

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US-08-936-772-3

US-08-936-772-3

US-09-395-918-3

US-09-639-881-14

US-08-660-542-21

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US-08-679-493A-156

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US-08-836-922-4

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US-09-639-881-3

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US-09-639-881-3

US-09-639-881-2

US-08-936-387-13

US-09-141-833-2

US-09-141-833-5

US-09-141-833-5

US-09-141-833-5

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US-08-936-387-16

US-08-936-387-18

US-08-936-387-18
                                                                                                                                                                                                                 US-08-347-492B-12

US-08-347-492B-12

Sequence 12, Application US/08347492B

Patent No. 5602008

GENERAL INFORMATION:
APPLICANT: Wilde, Craig G.
APPLICANT: Hawkins, Phillip R.
APPLICANT: Bandman, Olga
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: EXPRESSED CHEMOKINES, THI
TITLE OF INVENTION: PRODUCTION AND USES
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
CITY: Palo Alto
STATE: CA
COUNTRY: U.S.
                                                                                                                                                                                  ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE.

COMPUTER: IBM COMPATIBLE

COMPUTER: DOS

SOFTWARE: FastSEQ Version 1.5

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/347,492B

FILING DATE: 29-NOV-1994

PRICATION NUMBER: 08/303,241

FILING DATE: 07-SEP-1994

APPLICATION NUMBER: 08/320,011

FILING DATE: 05-OCT-1994

ATTORNEY/AGENT INFORMATION:

NAME: Luther, Barbara J

REGISTRATION NUMBER: PF-0024

TELECOMMUNICATION INFORMATION:

TELEPHONE: 415-855-0555

TELEFAX: 415-852-0195

INFORMATION FOR SEQ ID NO: 12:

SEQUENCE CHARACTERISTICS:

LENGTH: 91 amino acids
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MEDIUM TYPE: Diskette
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TOPOLOGY: li
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Sequence 5, Application US/08375346A
Patent No. 5605817
GENERAL INFORMATION:
APPLICANT: Coleman, Roger
APPLICANT: Wilde, Craig G.
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: A NEW CHEMOKINE EXPRESSED IN FETAL SPL
TITLE OF INVENTION: ITS PRODUCTION AND USES
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
STREET: 3330 HILLVIEW AVENUE
CITY: PALO ALTO
STATE: CALIFORNIA
COUNTRY: USA
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Score 478; DB 1;
Pred. No. 1e-47;
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Pred. No.
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REFERENCE/DOCKET NUMBER: PF-0026
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 855-0555
TELEFAX: (415) 855-0572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/375,34
FILING DATE: 19-JAN-1995
CLASSIFICATION: 435
PRIOR APPLICATION: 435
PRICATION NUMBER:
FILING DATE:
APPLICATION NUMBER:
FILING DATE:
APPLICATION NUMBER:
FILING DATE:
APPLICATION NUMBER:
APPLICATION NUMBER:
ATTORNEY/AGENT INFORMATION:
NAME: LUTHER, BARBARA J.
REGISTRATION NUMBER: 33,954
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th 100.0%; S Similarity 100.0%; P 91; Conservative 0;
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SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
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Pred. No. 1e-47;
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Mismatches
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TITLE OF INVENTION: NOVEL RANTES HOMOLOG
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc
STREET: 3174 Porter Drive
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 AVVFVTRKNROVCANPEKKWVREYINSLEMS
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APPLICANT: BANDMAN, OLGA; APPLICANT: COLEMAN, ROGER; APPLICANT: STUART, SUSAN G.
TITLE OF INVENTION: NEW CHEMOKINE; NUMBER OF SEQUENCES: 9; CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/633,68
FILING DATE: Filed Herewith
ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0063
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFHONE: 415-855-0195
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
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100.0%; Pr
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                                                                                                     3, Application US/08633682
5. 5840544
                                                                                                                                              Hawkins, Phillip R.
Bandman, Olga
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IBM Compatible
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: si
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                                                                                                                       Patent No. 5840544
GENERAL INFORMATION:
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IMMEDIATE SOURCE:
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APPLICANT:
APPLICANT:
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                                                                                          US-08-633-682-3
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Best Local
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Matches
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SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/421,144A
FILING DATE: 13-APR-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33954
REGISTRATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
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Pred. No. 1e-47;
Mismatches
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1-08-798-143-12
Sequence 12, Application US/08798143
Patent No. 5936068
GENERAL INFORMATION:
APPLICANT: Wilde, Craig G.
APPLICANT: Bandman, Olga
APPLICANT: Bandman, Olga
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: EXPRESSED CHEMOKINES, THI
TITLE OF INVENTION: PRODUCTION AND USES
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
CITY: Palo Alto
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/798,143
FILING DATE: 10-FEB-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
ADDRESSEE: INCYTE PHARMACEUTICALS, STREET: 3174 Porter Drive CITY: Palo Alto STATE: CA COUNTRY: USA ZIP: 94304
                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
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Similarity 100.0%;
91; Conservative (
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STRANDEDNESS: sing
TOPOLOGY: linear
MOLECULE TYPE: prote
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Best Local S
Matches 91
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APPLICANT: Wilde, Craig C.
APPLICANT: Wilde, Craig C.
APPLICANT: Seilhamer, Jeffrey J.
IITLE OF INVENTION: CHEMOKINE EXPRESSED IN FETAL SPLEEN,
ITTLE OF INVENTION: ITS PRODUCTION AND USES
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
CITY: Palo Alto
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Pred. No. 1e-47;
Mismatches
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,123B
FILING DATE: 06-JUN-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/375,346
FILING DATE: 19-JAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0026-1 DIV
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 AVVFVTRKNRQVCANPEKKWVREYINSLEMS 91
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APPLICATION NUMBER: 08/347,492
FILING DATE: 29-NOV-1994
APPLICATION NUMBER: 08/303,241
FILING DATE: 07-SEP-1994
APPLICATION NUMBER: 08/320,011
FILING DATE: 05-OCT-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                   PF-0024
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tert No. 5845506
ENERAL INFORMATION:
APPLICANT: Coleman, Roger
APPLICANT: Wilde, Craig C.
APPLICANT: Seilhamer, Jeffrey J.
                                                                                                                          NAME: Luther, Barbara J
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
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1; Conservative
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MEDIUM TYPE: Diskette
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IMMEDIATE SOURCE:
LIBRARY: GENBANK
CLONE: GI 134510
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SOFTWARE: FastSE
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STRANDEDNESS: si
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US-08-467-1238-5
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US-08-798-143-12
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Patent No
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APPLICANT: Hawkins, Phillip R.

APPLICANT: Bandman, Olga
APPLICANT: Murry, Lynn E.

TITLE OF INVENTION: NOVEL RANTES HOMOLOG FROM PROSTATE
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
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red. No. 1e-
Mismatches
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Pred. No.
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
ODERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/936,772
FILING DATE:
APPLICATION NUMBER: 08/633,682
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0063 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
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Pred.
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Sequence 3, Application US/08936772
Patent No. 6015883
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TELEFAX: 415-555-0555
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino 2017
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IMMEDIATE SOURCE:
LIBRARY: GenBank
CLONE: GI 134510
                                                                                                  single
                                                                                                                          peptide
                                                                                  TYPE: amino acid
STRANDEDNESS: sing
TOPOLOGY: linear
MOLECULE TYPE: pepti
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Best Local Similarity
Matches 91; Conser
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TELEFAX: 41
INFORMATION FOR
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                                 TELEFAX: 4: INFORMATION FOR
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Sequence 14, Application US/08836922

Patent No. 6159711

GENERAL INFORMATION:
APPLICANT: WELLS, TIMOTHY NIGEL CARL
APPLICANT: WELLS, TIMOTHY NIGEL CARL
TITLE OF INVENTION: COMPOSITIONS COMPRISING IT FOR TREATMENT OF INFLAMMATION
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
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  Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Batentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/836,922
FILING DATE: 23-MAY-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: GB 9424835.8
FILING DATE: 08-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9512319.6
FILING DATE: 16-JUN-1995
ATTORNEY/AGENT INFORMATION:
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100.0%; Pred. No. 1
cive 0; Mismatche
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REGISTRATION NUMBER: 32,955
REFERENCE/DOCKET NUMBER: 1430-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 816-4011
TELEFAX: (703) 816-4011
TELEFAX: (703) 816-4100
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acids
TOPOLOGY: linear
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1100 NORTH GLEBE ROAD
ARLINGTON
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       Conservative
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91; Conser
                                                                                                                                                                                                                                                                                                                                                                                                               VIRGINIA
                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A. ZIP: 22201-4714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
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                                                                                                                                                                                    RESULT 8
US-08-836-922-14
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US-09-395-918-3
       91;
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CITY: P
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                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/639,881
FILING DATE: 17-AUG-2000
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/836,922
FILING DATE: 23-MAY-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9424835.8
FILING DATE: 08-DEC-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                    SEE: NIXON & VANDERHYE P.C. 1100 NORTH GLEBE ROAD ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1430-163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 16-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: WILSON MARY:
                                                                                                                                                                                                                                                                                        : Floppy disk
IBM PC compatible
YSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21, Application US/08480449
Patent No. 5688927
GENERAL INFORMATION:
APPLICANT: Godiska, Ronald
APPLICANT: Gray, Patrick W.
            Sequence 14, Application US/09639881
Patent No. 6555105
GENERAL INFORMATION:
APPLICANT: INNES PROUDFOOT, AMAN
APPLICANT: WELLS, TIMOTHY NIGEL
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TELEPHONE: (703) 816-4011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         32,955
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TELEFAX: (703) 816-4100
INFORMATION FOR SEQ ID NO: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: WILSON, MARY J.
REGISTRATION NUMBER: 32,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          91 amino acids
amino acid
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                                                                                                                                  NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                    STATE: VIRGINIA COUNTRY: U.S.A. ZIP: 22201-4714
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                                                                                                                                                                                                                                                                         COMPUTER READABLE MEDIUM TYPE: F1
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US-09-639-881-14
                                                                                                                                                                      ADDRESSEE:
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US-08-480-449-21
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US-09-639-881-14
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STATE:
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APPLICANT: Chang, Yuan
APPLICANT: Bohenzky, Roy A
APPLICANT: Bohenzky, Roy B
TITLE OF INVENTION: USES THEREOF
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: 45185-G-PCT-US
CURRENT APPLICATION NUMBER: US/09/230,371A
CURRENT FILING DATE: 1999-11-17
PRICH APPLICATION NUMBER: PCT/US97/13346
PRICH FILING DATE: 1997-07-22
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red. No. 1e-47;
Mismatches 0
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100.0%; Pred. No. 1e-47;
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                                              APPLICANT: Hawkins, Phillip R.
APPLICANT: Bandman, Olga
APPLICANT: Murry, Lynn E.
TITLE OF INVENTION: DNA ENCODING RANTES HON
FILE REFERENCE: PF-0063 US
CURRENT APPLICATION NUMBER: US/09/395,918
CURRENT FILING DATE: 1999-09-14
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PERL Program
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61 AVVFVTRKNRQVCANPEKKWVREYINSLEMS
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Pred.
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 3, Application US/0939591
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Patent No. 6348586
GENERAL INFORMATION:
APPLICANT: Chang, Yuan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OF SEQ ID NOS:
                                                                                                                                                                                                                                                                        FEATURE:
NAME/KEY: misc feature
OTHER INFORMATION: Gen
3-09-395-918-3
                                                                                                                                                                                                                                                         ORGANISM: Homo sapiens
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   Sequence 3, Applicat
Patent No. 6238666
GENERAL INFORMATION:
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S-09-230-371A-25
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NO 25

NUMBER

Query Match Best Local S Matches 91

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APPLICANT: INNES PROUDFOOT, AMANDA ELIZABETH
APPLICANT: WELLS, TIMOTHY NIGEL CARL
TITLE OF INVENTION: RANTES PEPTIDE AND FRAGMENTS AND
TITLE OF INVENTION: COMPOSITIONS COMPRISING IT FOR TREATMENT OF INFLAMMATION
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ower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Godiska, Ronald
APPLICANT: Gray, Patrick W.
TITLE OF INVENTION: MACROPHAGE DERIVED CHEMOKINE
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall, O'Toole, Gerstein, Murray
STREET: 6300 Sears Tower, 233 South Wacker Driv
                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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S-08-660-542-21
Sequence 21, Application US/08660542
Patent No. 5932703
GENERAL INFORMATION:
APPLICANT: Godiska, Ronald
APPLICANT: Gray, Patrick W.
TITLE OF INVENTION: MACROPHAGE DERIVED CHEMOKINE AND CHE
TITLE OF INVENTION: ANALOGS
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & Borun
STREET: 6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Borun
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STATE: Illinois
COUNTRY: United States of America
ZIP: 60606-6402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/660,542
                                                                                                                                                  #1.25
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CITY: Chicago

STATE: Illinois

COUNTRY: United States of America

ZIP: 60606-6402

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: BATCATION DATA:

APPLICATION NUMBER: US/08/480,449

FILING DATE:

CLASSIFICATION: 530

ATTORNEY/AGENT INFORMATION:

NAME: Gass, David A.

REGISTRATION NUMBER: 38,153

REGISTRATION INFORMATION:

TELEPHONE: 312/474-6300

TELEPAX: 312/474-6448

TELEFAX: 25-3856

INFORMATION FOR SEQ ID NO: 21:

SEQUENCE CHARACTERISTICS:

LENGTH: 91 amino acids

TYPE: amino acid
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Pred. No. 5.1e-47;
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N: "RANTES
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CLASSIFICATION: 514
IOR APPLICATION DATA:
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OTHER INFORMATION:
3-08-480-449-21
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Pred. No. 5.1e-47;
0; Mismatches 1;
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No. 5.1e-47;
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US-08-679-493A-155
Sequence 155, Application US/08679493A
Patent No. 6303295
GENERAL INFORMATION:
TITLE OF INVENTION:
FILE REFERENCE: 55-95
CURRENT APPLICATION NUMBER: US/08/679,493A
CURRENT FILING DATE: 1996-07-12
PRIOR APPLICATION NUMBER: 60/001203
PRIOR FILING DATE: 1995-07-14
PRIOR FILING DATE: 1995-07-14
PRIOR FILING DATE: 1995-09-01
NUMBER OF SEQ ID NOS: 216
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 155
LENGTH: 91
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 472;
Pred. No. 5
                                                                                                  38,153
ER: 27866/33318
                                         APPLICATION NUMBER: 08/479,620
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Gass, David A.
REGISTRATION NUMBER: 38,153
REFERENCE/DOCKET NUMBER: 27866/3:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/474-6300
TELEFAX: 312/474-648
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 21:
FILING DATE: 16-NOV-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/479 620
FILING DATE:
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ilarity 98.9%;
Conservative (
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larity 98.9%;
Conservative
                                                                                                                                                                                                      LENGTH: 91 amino acids
TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                        feature
                                                                                                                                                                                                                                                             peptide
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US-08-679-493A-155
                                                                                                                                                                                                                                                                                                                                Protein
1..68
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90; Conserv
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LOCATION:
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SULT 15
-08-479-603-21
Sequence 21, Application US/08479603
Patent No. 6320023
GENERAL INFORMATION:
APPLICANT: Godiska, Ronald
APPLICANT: Godiska, Rarshall, O'Toole, Gerstein, Murray & BONTSTEET 6300 Sears Tower, 233 South Wacker Drive
STREET: 6300 Sears Tower, 233 South Wacker Drive
CITY: Chicago
STREE: Illinois
COUNTRY: United States of America
ZIP: 6060-6402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: BAPENCH Release #1.0, Version #1.25
COMPUTER: BAPENCH NO BATA:
APPLICATION: S30
CURRENT APPLICATION: S30
ATTORNEY/AGENT INFORMATION:
NAME: Gass, David A:
REGISTRATION TOFORMATION:
NAME: Gass, David A:
REGISTRATION TOFORMATION:
TELEPHONE: 312/474-6300
TELEPHONE: 312/474-6300
TELEFRAX: 25-3856
INFORMATION FOR REAL STICS:
LENGTH: 91 amino acids
TYPE: amino acids
TTYPE: amino acids
TTYPE: mino acids
TTYPE: MOLECTYPE: peptide
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OTHER INFORMATION: "RANTES"
:-08-479-603-21
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GenCore version 5.1.6
(c) 1993 - 2003 Compugen Ltd
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using sw model protein - protein search, 2003, 15:36:20 December 16

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SUMMARIES

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AjRocession: A31767
AjMolecule type: mRNA
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AjRocession: A31767
AjMolecule type: mRNA
AjRocession: B30574
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AjRocession: B30574
AjMolecule type: mRNA
AjRocession: A3741; MUID: 89325421; PMID:2568930
AjTitle: Isolation and characterization of a cDNA encoding a putative cytokine which i AjRocession: A37411; MUID: 89325421; PMID:2568930
AjRocession: A37411; MUID: 8932035; PIDN: CAA34291.1; PID:g32036
AjRocession: B30574
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RESULT : A31767 macroph?	in i	ato	je č	ein HC21	ALIGNMENT in 1-beta precui	Cu Line] - human g; lymphocyte activation ge
ជា-ក ២០៩២៦១១	ies: Homo sies: Homo sies: 07-Jun-19 ssion: JH03 eras, E.; R mmunol. 27, e: Cloning erence numberssion: JH03	1 T- 19; # 19; # 10; 10; 10; 10; 10; 10; 10; 10; 10; 10;	(man (man quenc 0978; 0man, 1102, 1102, 319;		ion prision 67; A3 Jitsuk	gamma 7-1998 #text B30574; B458 S.; Genevee, yte activatio MID:2247088	change 15-Sep-2000 17; D30552 C.; Mechiche, S.; Viegas-Pequ n gene (LAG-1).
A; Status A; Molecy A; Resid A; Exper R; Napol A; Title A; Refer A; Acces	ttus: translar ecule type: 1 idues: 1-92 desimental son berimental son bolitano, M.; ol. Chem. 26 iol. Chem. 26 iol. The gene erence numbers	ONA ONA ABAI> B: GB LICE: Modi 6. 17 enco	x5368 x5368 natur w.S. 31-17	G G G G G	Printer Control of Con); PIDN:CAA37723.1 1, strain CD3-CD2- J.; Gnarra, J.R.; kine. Genomic stru ; PMID:1894635	3.1; PID:934218 CD2+, F5, 5111E5 R.; Seuanez, H.N.; Leonard, W.J Btructure, HTLV-I/tax responsiv
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A; Forsidues: 1-92 camp.
A; Fresidues: 1-92 camp.
A; Freference camp.
A; Fresidues: 1-92 camp.
A; Accession: JL0088
A; MUD: 8966; PIDN: AAA39708.1; PID: 99697
A; Accession: P80304
A; Molecule type: protein
A; Fresidues: 1-92 camp.
A; Accession: P80304
A; Molecule type: protein
A; Fresidues: 24-33, 'XX', 38 camp.
A; Accession: P80304
A; Molecule type: protein
A; Fresidues: 24-33, 'XX', 38 camp.
A; Accession: P80304
A; Molecule type: protein
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                                                                                                            macrophage inflammatory protein 1-beta precursor - mouse
N;Alternate names: H400; SIS gamma; T-cell activation protein gamma
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 28-Aug-1989 #sequence revision 28-Aug-1989 #text_change 16-Jul-1999
C;Accession: C30552; JL0088; PS0304; S22042
C;Accession: C30552; JL0088; PS0304; S22042
R;Brown, K.D.; Zurawski, S.M.; Mosmann, T.R.; Zurawski, G.
J. Immunol. 142, 679-687, 1989
A;Title: A family of small inducible proteins secreted by leukocytes are members of various activation processes.
A;Reference number: A30552; MUID:89093958; PMID:2521353
A;Accession: C30552
A;Accession: C30552
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C; Species: Mus musculus (house mouse)

C; Species: Mus musculus (house mouse)

C; Species: Mus musculus (house mouse)

C; Species: 17-Jul-1992 #sequence revision 17-Jul-1992 #text change 16-Jul-1999

C; Accession: S11685; A32393; S04533; A53885; A30552; PS0303; A27596; I56104

R; Grove, M.; Lowe, S.; Graham, G.; Pragnell, I.; Plumb, M.

Nucleic Acids Res. 18, 5561, 1990

A; Title: Sequence of the murine haemopoietic stem cell inhibitor/macrophage
A; Reference number: S11685; MUID: 91016858; PMID: 2216738

A; Accession: S11685
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larity 76.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Superfamily: macrophage inflammatory protein; Superfamily: macrophage inflammation; Keywords: chemotaxis; Cytokine; inflammation; 1-23/Domain: signal sequence #status predicted <SIG>; 24-92/Product: macrophage inflammatory protein 1-beta #status experimen; 34-58, 35-74/Disulfide bonds: #status experimental
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%; Residues: 1-92 < MOR>
%; Cross-references: GB:D17402; NID:g599577; PIDN:BAA04226.1; PID:g599578
5; Superfamily: macrophage inflammatory protein
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mmune activation gene 2 - rabbit

;;Species: Oryctolagus cuniculus (domestic rabbit)

;;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change 16-Jul-

;;Accession: 146730

;;Mori, S.; Goto, K.; Goto, F.; Mutakami, K.; Ohkawara, S.; Yoshinaga,

;mori, S.; Goto, K.; Goto, F.; Mutakami, R.; Ohkawara, S.; Yoshinaga,

;nt. Immunol. 6, 149-156, 1994

;rtile: Dynamic changes in mRNA expression of neutrophils during the call the call
                                          lymphocytes
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                                                                                                                                                                                                                           ; PIDN:AAA36752.1;
R.; Zurawski, G.
                                          H
                                      activated human
; PMID:2809212
                                                                                                                                                                                                                                      Brown, K.D.; Zurawski, S.M.; Mosmann, T.R.; Zurawski, Immunol. 142, 679-687, 1989
Title: A family of small inducible proteins secreted of various activation processes.
Reference number: A30552; MUID:89093958; PMID:2521353; Accession: D30552
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Pred. No. 2.3e-45;
Mismatches 0;
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red. No. 4.6e-37;
Mismatches 7
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preliminary; translated from GB/EMBL/DDBJ
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Title: A novel polypeptide secreted by activ Reference number: A45817; MUID:90038522; PMI Accession: B45817

Molecule type: mRNA
Residues: 7-55,'I',57-79,'T',81-92 <MIL>
Cross-references: GB:M57503; NID:9339726; PI Brown, K.D.; Zurawski, S.M.; Mosmann, T.R.;
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Pred. No. 4
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Cross-references: GDB:127451; OMIM:153335
Map position: 17q21-17q21
Introns: 26/1; 64/2
Superfamily: macrophage inflammatory protestions: Superfamily: macrophage inflammatory protestions
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Similarity 82.6%;
76; Conservative
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1; Residues: 1-92 < MOR>
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Best Local
Matches 9
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                                                                                                                                                                                                     60 PGVIFLTKRNRQICADSKETWVQEYITDLELN 91
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 A; Introns: 26/1; 64/2
 C; Superfamily: macrophage inflammatory protein C; Superfamily: macrophage inflammatory protein C; Keywords: cytokine C; Keywords: cytokine F; 1-22/Domain: signal sequence #status predict F; 1-22/Domain: signal sequence #status predict
                                                                                                                                                                  PAVVFOTKRSKOVCADPSESWVQEYVYDLELN
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llarity 58.7%;
Conservative 20
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Molecule type: mRNA
Residues: 1-92 - cDAV.
Cross-references: EMBL:X12531; NID:g53122; PIDN:CAA31047.1; PID:g53123
Brown, K.D.; Zurawski, S.M.; Mosmann, T.R.; Zurawski, G.
Immunol. 142, 679-687, 1389
Inmunol. 142, 679-687, 1389
Inmunol. 142, 679-687, 1389
Inmunol. 142, 679-687, 1389
Infile: A family of small inducible proteins secreted by leukocytes are members of a ne of various activation processes.
Reference number: A30552; MUID:89093958; PMID:2521353
Accession: A30552
Molecule type: mRNA
Residues: 1-21, L', 23-61, 47, 63-92 - cBRO>
Residues: 1-21, L', 23-61, 47, 80-92 - cBRO>
Residues: 1-21, L', 23-61, 80-98 - components of macrophage inflammatory protein 1, and clon sherry. Bs. Tekamp-Olson, P.; Gallegos, C.; Bauer, D.; Davatelis, G.; Sherry, B.; Reference number: JL0088; MUID:89067830; PMID:3058856
Accession: PS0303
Residues: 24-33, XXX, 36-54 - cSHE>
ROBIOLE type: mRNA
Residues: 24-33, XXX, 36-54 - cSHE>
Residues: 25-581, 1988
Residues: 25-591, 1988
Residues: 1-21, 1988
Residue
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Cross-references: GB:J04491; NID:g201524; PIDN:AAA40304.1; PID:g201529
Cross-references: GB:J04491; NID:g201524; PIDN:AAA40304.1; PID:g201529
Davatelis, G.; Tekamp-Olson, P.; Wolpe, S.D.; Hermsen, K.; Luedke, C.
Exp. Med. 167, 1939-1944, 1988
Title: Cloning and characterization of a cDNA for murine macrophage in Reference number: S04533; MUID:88258380; PMID:3290382
Accession: S04533
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Residues: 1-92 <GRO>
Cross-references: EMBL:X53372; NID:g54062; PIDN:CAA37452.1; PID:g2979
Cross-references: EMBL:X53372; NID:g54062; PIDN:CAA37452.1; PID:g2979
Note: the authors' translation of the nucleotide sequence differs at Kwon, B.S.; Weissman, S.M.
Co. Natl. Acad. Sci. U.S.A. 86, 1963-1967, 1989
Title: cDNA sequence of two inducible T-cell genes.
Reference number: A32393; MUID:89184547; PMID:2784565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Residues: 1-48, 'E', 50-90, 'I', 92 <DA2>
Cross-references: EMBL:X12531
Note: the authors translated the codon GAG for residue 49 as Asp ar Note: the sequence has been corrected in reference A53885
Davatelis, G.; Tekamp-Olson, P.; Wolpe, S.D.; Hermsen, K.; Luedke, Exp. Med. 170, 2189, 1989
Reference number: A53885
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This protein is a monokine.
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Similarity 60.9%; Pred. No. 1.6e-27;
56; Conservative 16; Mismatches 19;
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Superfamily: macrophage inflammatory protein
Keywords: heparin binding
1-23/Domain: signal sequence #status predicted
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Best Local S
Matches 56
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Balsey, and protein precursor - human NyAlternate names: macrophage inflammatory protein homolog GOS19-2; small inducible cy Cispecies: Homos sapiens (man) Cispecies: Homos sapiens (man) Cipate: 28-Sep-1990 #sequence revision 28-Sep-1990 #text_change 20-Jun-2000 C;Accession: B15673; B10412; S10157; B30908 Rimada, K. Mol. Cell. Biol. 10, 3646-3658, 1990 A;Title: Structures of human genes coding for cytokine LD78 and their expression. A;Reference number: A35673; MUID:90287155; PMID:1694014 A;Accession: B35673 MUID:90287155; PMID:1694014 A;Accession: B35673 MUID:90287155; PMID:1694014 A;Accession: B35673 MUID:90287155; PMID:1694014 A;Accession: B3673 MUID:90145; NID:219907; PIDN:BAA14173.1; PID:9219908 A;Accession: B369-602, 1990 A;Accession: B369-602, 1990 A;Attle: Three human homologs of a murine gene encoding an inhibitor of stem cell prol A;Accession: B30412 MUID:91103879; PMID:2271120 A;Accession: B30412 MUID:91103879; PMID:2271120 A;Accession: B30412 MUID:91103879; PMID:2271120 A;Accession: B30412 MUID:91103879; MUID:9182848; PIDN:AAA35889.1; PID:9182849 A;Accession: B30412 MUID:910404 MUID:90287702; PMID:1972563
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Comment: This protein is a member of a "small inducible" or "activation specific"
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 29-May-1998 #sequence_revision 29-May-1998 #text_change 16-Jul-1999
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:: LD78-beta protein #status predicted
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Pred. No. 1.8e-27;
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monocyte chemoattractant cytokine RANTES precursor - human
Nighternate names: small inducible cytokine A5; T-cell specific cytokine RANTES
C;Species: Homo sapiens (man)
C;Scossion: A28815
A;Title: Aluman T cell-specific molecule is a member of a new gene family.
A;Title: A human T cell-specific molecule is a member of a new gene family.
A;Molecule type: mRNA
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                                                                                                                                               1 MKLCVTVLSLLMLVAAFCSPALSAPMGSDPPTACCFSYTARKLPRNFVVDYYETSSLCSQ
                                                                                                                                                                             monocyte adherence-induced protein 5 beta - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 06-Nov-1992 #sequence_revision 06-Nov-1992 #text_cha
C;Accession: C60407
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                                              61.8%; Score 300.5; DB 2; larity 57.6%; Pred. No. 2.3e-25; Conservative 19; Mismatches 19;
                                                                                                                                                                                                                                                            PAVVFOTKRSKOVCADPSESWVQEYVYDLELN 92
F;33-57,34-73/Disulfide bonds: #status predicted
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Species: Homo sapiens (man)
Date: 03-Aug-1992 #sequence revision 03-Aug-1992 #text_change 21-
Date: 03-Aug-1992 #sequence revision 03-Aug-1992 #text_change 21-
Accession: A35673; A30574; A30412; A24198; A30908
Nakao, M.; Nomiyama, H.; Shimada, K.

J. Cell. Biol. 10, 3646-3658, 1990
Title: Structures of human genes coding for cytokine LD78 and the Reference number: A35673; MUID:90287155; PMID:1694014
Reference number: A35673; MUID:90287155; PMID:1694014
References: GB:D90144; NID:g219905; PIDN:BAA14172.1; PID:g2
Cross-references: GB:D90144; NID:g219905; PIDN:BAA14172.1; PID:g2
Cross-references: GB:D90144; NID:g219905; PIDN:BAA14172.1; PID:g2
Title: Mitogenic activation of human T cells induces two closely
Reference number: A30574; MUID:89140347; PMID:2521882
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names: LD78-alpha protein precursor; lymphocyte tumor
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                                      Shi, M.M.; Godleski, J.J.; Paulauskis, J.D.
Shi, M.M.; Godleski, J.J.; Paulauskis, J.D.
ochem. Biophys. Res. Commun. 211, 289-295, 1995
Title: Molecular cloning and posttranscriptional regulation
Reference number: I52322; MUID:95298037; PMID:7779098
Accession: I52322
Status: preliminary; translated from GB/EMBL/DDBJ
Molecule type: mRNA
Residues: 1-92 <RES>
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;Cross-references: GB:M23178; GB:M32337; NID:g182846; PIDN:
;Obaru, K.; Fukuda, M.; Maeda, S.; Shimada, K.
Biochem. 99, 885-894, 1986
;Title: A cDNA clone used to study mRNA inducible in human
;Reference number: A24198; MUID:86223879; PMID:3086300
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Similarity 57.6%; Pred. No. 1.1e-25;
53; Conservative 17; Mismatches 21;
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; Cross-references: GDB:120368; OMIM:182283
; Map position: 17q11-17q21
; Superfamily: macrophage inflammatory protein
; 1-20/Domain: signal sequence #status predicted
; 21-92/Product: macrophage inflammatory protein
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Cross-references: EMBL:U22414; NID:g790632; 3
Superfamily: macrophage inflammatory protein
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Shi, M.M., Godles
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monocytes but not neutrophils is an :
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Reference number: A46539; MUID:92z830000, ....

Accession: A46539

Accession: A46539

Molecule type: mRNA

Molecule type: mRNA

Residues: 1-18,'A',20-91 <SCH>

Residues: 1-18,'A',20-91 <SCH

Residues: 1-18,'A',20-91 <SC
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words: chemotaxis; cytokine; immediate-early protein; inflammation
3/Domain: signal sequence #status predicted <SIG>
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Neilson, E.G.; Krensky, A.
dney Int. 41, 220-225, 1992
Title: Isolation and characterization of cDNA from renal tubular
Reference number: I56970; MUID:92277990; PMID:1375672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cytokine RANTES #status
                                                                                                                                                                                                                                                                                                                                                                                                         P.S.; Neilson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 91,
                                                                                                                                                                                                                                                                                                                                    change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reference number:
Accession: 148875
Status: preliminary; translated from GB/EMBL/DDBJ
Molecule type: DNA
Residues: 1-91 <DAN>
Cross-references: EMBL:U02298; NID:g460090; PIDN:AAA18302.1;
Cross-references: EMBL:U02298; NJY.
Cross-references: EMBL:U02298; NJY.
Schall, T.J.; Simpson, N.J.; Mak, J.Y.
r. J. Immunol. 22, 1477-1481, 1992
Title: Molecular cloning and expression of the murine RANTES
Peference number: A46539; MUID:92289805; PMID:1376260
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 element
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Status: translated from GB/EMBL/DDBJ
Molecule type: mRNA
Molecule type: mRNA
Residues: 1-40,'E',42-91 <NEI>
Cross-references: GB:M77747; NID:g200649; PIDN:AAA40029.1; Comment: This chemoattractant for monocytes but not neutropl Genetics:
Introns: 26/1; 63/2
Superfamily: macrophage inflammatory protein
Keywords: chemotaxis; cytokine; immediate-early protein; in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           translation not shown; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                               mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Note: sequence extracted from NCBI backbone (NCBIN:106 Shin, H.S.; Drysdale, B.E.; Shin, M.L.; Noble, P.W.; I. Cell. Biol. 14, 2914-2925, 1994
Title: Definition of a lipopolysaccharide-responsive Reference number: I48654; MUID:94217689; PMID:7513046 Accession: I48654
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           enomic organization, and chromosomal 148875; MUID:94132613; PMID:7507961
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                                                                                                                                                                                                                      nocyte chemoattractant cytokine RANTES precursor Alternate names: MuRantes Species: Mus musculus (house mouse)
Date: 18-Jun-1993 #sequence revision 16-Aug-1996
Accession: 148875; A46539; 148654; 156970
Danoff, T.M.; Lalley, P.A.; Chang, Y.S.; Heeger, Immunol. 152, 1182-1189, 1994
Title: Cloning, genomic organization, and chromos Reference number: 148875; MUID:94132613; PMID:750
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4e-19;
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Pred. No. 4e-
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45; Conser
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Molecule type: I
Residues: 1-91
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24-91/Product:
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NyAlternate names: monocyte chemoattractant protein MCP-3
C;Species: Homo sapiens (man)
C;Species: Homo sapiens
C;Accession: A54678; JC1478; 532222
C;Accession: A54678; JC1478; 532222
R;Opdenakker, G; Fiten, P.; Nys, G; Froyen, G; Van Roy, N.; Speleman, F.; Laureys, Genomics 21, 403-408, 1994
A;Title: The human MCP-3 gene (SCYA7): cloning, sequence analysis, and assignment to the A;Reference number: A54678
A;Moclecule type: DNA
A;Residues: 1-109 <OPD>
A;Cross-references: GB:X2309
B;Opdenakker, G; Froyen, G; Fiten, P.; Proost, P.; Van Damme, J.
A;Residues: 1-109 <OPD>
A;Residu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Accession: S32222
A;Accession: S32222
A;Molecule type: mRNA
A;Residues: 1-109 <MIN>
A;Cross-references: EMBL:X71087; NID:g288396; PIDN:CAA50405.1; PID:g288397
A;Cross-references: EMBL:X71087; NID:g288396; PIDN:CAA50405.1; PID:g288397
C;Comment: This protein induces proteinase secretion and chemotaxis by macrophages and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   a human monocyte-derived monocyte chemoattr
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N.Alternate names: monocyte chemoattractant protein-1
C.Species: Rattus norvegicus (Norway rat)
C.Species: Rattus norvegicus (Norway rat)
C.Battus norvegicus (Norway rat)
C.Accession: S07723; UN0128
R.YIMmers, H.T.M.; Pronk, G.J.; Bos, J.L.; van der Eb, A.J.
Nucleic Acids Res. 18, 23-34, 1990
A.Title: Analysis of the rat UE gene promoter identifies an AP-1 binding site essentians 507723; MUID:90174947; PMID:2106664
A.Reference number: S07723; MUID:90174947; PMID:2106664
A.Residues: 1-148 «TIM»
A.Residues: 1-148 «TOS»
A.MOICOLE NUMBER: MUID:91128376; PMID:1704226
A.Accession: UN0128
A.MOICOLE COMMUNIANA
A.Residues: 1-148 «YOS»
A.Residues: 1-148 «Y
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Pred. No. 4.6e-14;
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larity 41.9%;
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monocyte chemoattractant protein-1 precursor - pig
C;Species: Sus scrofa domestica (domestic pig)
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 30-Sep-1993 #sequence_revision 20-Aug-1994 #text_change 16-Jul-1999
C;Date: 30-Sep-1993 #sequence_revision 20-Aug-1994 #text_change 16-Jul-1999
C;Date: 30-Sep-1993 #sequence_revision 20-Aug-1994 #text_change 16-Jul-1999
C;Date: Dorotone Unit of States (Size of Mana)
R;HOsang, K.; Knoke, I.; Klaudiny, J.; Wempe, F.; Wuttke, W.; Scheit, K.H.
R;Accies or occine luteal cells express monocyte chemoattractant protein-1 (MCP-1): Anal
A;Reference number: JC2136; MUD:94183284; PMID:7510962
A;Residues: 1-99 eHOS>
A;Cross-references: GB:248479; NID:9683716; PIDN:CAA88370.1; PID:9683717
A;Residues: 1-99 eHOS>
A;Cross-reference number: S57497
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-99 eACO>
A;Cross-references: EMBL:X79416; NID:9972312; PIDN:CAA55945.1; PID:9872313
C;Superfamily: macrophage inflammatory protein
C;Keywords: glycoprotein
F;1-37Domain: signal sequence #status predicted eSiG>
F;24-99/Product: monocyte chemoattractant protein-1 #status predicted
F;94/Binding site: carbohydrate (Asn) (covalent) #status predicted
J. Immunol. 150, 5025-5032, 1993
A; Title: cDNA cloning of guinea pig monocyte chemoattractant protein-1 and expression A; Title: cDNA cloning of guinea pig monocyte chemoattractant protein-1 and expression: A; Reference number: 148147
A; Reference number: 148147
A; Status: preliminary; translated from GB/EMBL/DDBJ
A; Molecule type: mRNA
A; Molecule type: mRNA
A; Residues: 1-120 < RES>
A; Cross-references: GB:L04985; NID:g349820; PIDN:AAA37047.1; PID:g349821
C; Genetics:
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C; Superfamily: macrophage inflammatory protein
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35.2%; Pred. No. 2.1e-12;
ive 23; Mismatches 35
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;Cross-references: GB:M19681; NID:g193486; PIDN:AAA37684.1;
;Kawahara, R.S.; Deuel, T.F.
Biol. Chem. 264, 679-682, 1989
;Title: Platelet-derived growth factor-inducible gene JE is;
;Reference number: A44771; MUID:89093129; PMID:2910858
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Species: Mus musculus (house mouse)

Date: 10-Sep-1999 #sequence revision 10-Sep-1999 #text

Accession: A30209; A44771; A30861

Rollins, B.J.; Morrison, E.D.; Stiles, C.D.

roc. Natl. Acad. Sci. U.S.A. 85, 3738-3742, 1988

;Title: Cloning and expression of JE, a gene inducible

;Reference number: A30209; MUID:88234501; PMID:3287374

;Accession: A30209

;Molecule type: DNA
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148147
nonocyte chemoattractant protein-1 - guinea pig
2;Species: Cavia porcellus (guinea pig)
2;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996
2;Accession: 148147
2;Yoshimura, T.
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                                                                                                               Introns: 36/1, 75/2
Superfamily: macrophage inflammatory protein
Keywords: cytokine, glycoprotein; inflammation
1-33/Domain: signal sequence #status predicted
34-109/Product: monocyte chemotactic protein 3
39/Binding site: carbohydrate (Asn) (covalent)
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larity 38.5%; Pred. No. 1.5
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uperfamily: macrophage inflammat
eywords: cytokine; glycoprotein
26/Binding site: carbohydrate (A
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Gene: GDB:SCYA7; SCYA6; MCP-3
Cross-references: GDB:138473;
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GenCore version 5.1.6 (c) 1993 - 2003 Compugen Ltd.
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protein - protein search, using sw model

December 16, 2003, 15:36:20; Search time 10.5574 Seconds (without alignments) 409.804 Million cell updates/sec no u

US-09-920-137A-7 486 tle: rfect score:

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st-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_41:* tabase :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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MEDLINE=94182137; PubMed=8134838;

MEDLINE=94182137; PubMed=8134838;

Lodi P.J., Garrett D.S., Kuscewski J., Tsang M.L.S., Weatherbee J.A.,

Leonard W.J., Gronenborn A.M., Clore G.M.;

Leonard W.J., Gronenborn A.M., Clore G.
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Birren B., Fasman K., McKernan K., Nusbaum C., Richardson P.,
Lander E.,
Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases.

[8]
SEQUENCE OF 6-92 FROM N.A.
MEDLINE=90038522; PubMed=2809212;
MEDLINE=90038522; PubMed=2809212;
Miller M.D., Hata S., Waal Malefyt R., Krangel M.S.;
Miller M.D., Hata S., Waal Malefyt R., Trangel M.S.;
"A novel polypeptide secreted by activated human T lymphocytes.
J. Immunol. 143:2907-2916(1989).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and MIP-1 beta as T cells.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gallo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W., Spinetti
Santoni A.,
                                                                                           ic structure, chromosomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              macrophage
the CCR8 r
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INDUCTION: By mitogens.
SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY
C-C) (CHEMOKINE CC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PubMed=8525373;
A.L., Garzino-Demo A., Arya S.K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for
                                                          Gnarra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and
                                                                                                Genomic
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                                                                                                              and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Luini
                                                                                                                                                                                                                                                                                                                                                                                                               [9]
RECEPTOR INTERACTION.
MEDLINE=98180363; PubMed=9521068;
Bernardini G., Hedrick J., Sozzani S., Luini Bernardini G., Hedrick J., Sozzani S., Luini Weiss M., Menon S., Zlotnik A., Mantovani A. Napolitano M.;
"Identification of the CC chemokines TARC an protein-1 beta as novel functional ligands for Eur. J. Immunol. 28:582-588(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           n of the CC chemokines TARC as as novel functional ligands 1. 28:582-588(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               alpha, a
by CD8+
                                                                                                                   sednences,
                                                        s.J.,
                                                                                              cytokine
                                                                                                                                                          266:17531-17536(1991)
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                                         1894635;
Cevario
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lusso P.;
Lusso P.;
"Identification of RANTES, MIP-1
HIV-suppressive factors produced
crience 270:1811-1815(1995).
                                    PubMed=1894635
H W.S., Cevario
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ALT
                                                                                                                       upstream
                                                                                                 Act-2
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AAA57256.1;
CAA37723.1;
CAA37722.2;
CAA37722.2;
AAB00790.1;
AAB00790.1;
                                                                                                    the
[6]
SEQUENCE FROM N.A.
MEDLINE=91373378; Pub
Napolitano M., Modi W
                                                                                                                      Š
                                                                               .J.;
encoding
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Cocchi F., Devico
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M25316;
J04130;
X53683;
X53682;
X16166;
M69203;
M69201;
                                                                                                                                                ٠.
                                                                                                                      responsiveness
                                                                                                                                            localization."
                                                                                                                                                                Chem.
                                                                                                    gene
                                                                                  Leonard W
"The gene
                                                                                                                                                                Biol.
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() (ACT-2).
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                                                                                                                                                                        Genew; HGNC:1000-0.

MIN; 182284; -.

GO; GO:0005615; C:extracellular space; TAS.

GO; GO:0008009; F:chemokine activity; TAS.

GO; GO:0004716; F:receptor signaling protein tyrosine kinase.

GO; GO:0007155; P:cell adhesion; TAS.

GO; GO:0007267; P:cell andioxinativ; TAS.

GO; GO:0007267; P:cell motility; TAS.

GO; GO:0007267; P:cell motility; TAS.

GO; GO:0007163; P:establishment and/or maintenance of cell po.

GO; GO:0006955; P:immune response; TAS.

GO; GO:0006955; P:imflammatory response; TAS.

GO; GO:000615; P:response to viruses; TAS.

GO; GO:0006165; P:signal transduction; TAS.

GO; GO:0008166; P:viral replication; TAS.

InterPro; IPR001811; Chemokine_sml.

R Pfam; PF00048; ILB; 1.

R PROSITE; PS00472; SMALL CYTOKINES_CC; 1.

R SIMALL SMO0199; SCY; 1.

T SIGNAL

T SIGNAL

T CHAIN

CHAIN

EN SIMILARITY.
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SYO4 RABIT
STANDARD;
PRT;
92 AA.
P46632;
01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Small inducible cytokine A4 precursor (CCL4) (Macrophage protein 1-beta) (MIP-1-beta) (Immune activation protein CCL4 OR SCYA4.
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Pred. No. 6.1e-47;
Mismatches 0;
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(IN REF.
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etazoa: Chordata;
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74
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Conservative
                                                                                                          IHUM; 30-APR-94.
IHUN; 30-APR-94.
1JE4; 03-OCT-01.
w; HGNC:10630; CCL4.
                                                 ; -; NOT_ANI
AAA36752.1;
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EMBL; AC003976; -; h
EMBL; M57503; AAA367
PIR; JH0319; A31767.
PDB; 1HUM; 30-APP
PDB; 1HUN; 20-APP
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92; Conser
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Eukaryota; N
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DISULFID
CONFLICT
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SY04 I
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=Long Evans; TISSUE=Lung;
Jones M.L., Shanley T.P., Schmal H., Friedl H.P., Ward P.A.;
Submitted (FEB-1994) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: MONOKINE WITH INFLAMMATORY AND CHEMOKINETIC PROPERTIES
                                                                                                       SEQUENCE FROM N.A.
STRAIN=New Zealand white;
STRAIN=New Zealand white;
MEDLINE=94198229; PubMed=8148323;
Mori S., Goto K., Goto F., Mutakami K., Ohkawara S., Yoshinaga M.;
Mori S., Goto K., Goto F., Mutakami C., Ohkawara S., Yoshinaga M.;
"Dynamic changes in mRNA expression of neutrophils during the course of acute inflammation in rabbits.";
of acute inflammation in rabbits.";
Int. Immunol. 6:149-156(1994).
Int. Immunol. 6:149-156(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inflammator
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUBUNIT: Homodimer (By similarity).
SUBCELLULAR LOCATION: Secreted.
SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY
C-C) (CHEMOKINE CC).
    Oryctolagua
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CYTOKINE
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cis; Inflammatory response; Signal.
23 BY SIMILARITY.
92 SMALL INDUCIBLE CYTOKINE
58 BY SIMILARITY.
74 BY SIMILARITY.
10066 MW; ECBA8818D42A735C CRC64;
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Pred. No. 1.9e-38;
); Mismatches 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P50230;
01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Small inducible cytokine A4 precursor (CCL4)
protein 1-beta) (MIP-1-beta).
CCL4 OR SCYA4 OR MIP1B.
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PIR; I46730; I46730.

HSSP; P13236; 1HUM.

InterPro; IPR000827; CC_chemkine_sml.

InterPro; IPR001811; Chemokine_ILB.

Pfam; PF00048; ILB; 1.

SMART; SM00199; SCY; 1.

PROSITE; PS00472; SMALL_CYTOKINES_CC; Cytokine; Chemotaxis; Inflammatory resignate

SIGNAL

CHAIN

DISULFID

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92

BY SIMIL

DISULFID

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74

BY SIMIL
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       Lagomorpha;
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82.6%;
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    Mammalia; Eutheria;
NCBI_TaxID=9986;
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76; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (BY SIMILARITY
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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Brown K.D., Zurawski S.M., Mosmann T.R., Zurawski G.;
"A family of small inducible proteins secreted by leukocytes are
members of a new superfamily that includes leukocyte and fibroblast-
derived inflammatory agents, growth factors, and indicators of various
activation processes.";
J. Immunol. 142:679-687(1989).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (Macrophage inflammatory (SIS-gamma) (ACT2).
SUBUNIT: Homodimer (By similarity).
SUBCELLULAR LOCATION: Secreted.
SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY (SMALL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi
Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=89067830; PubMed=3058856; Sherry B., Tekamp-Olson P., Gallegos C., Bauer D., Davatelis G. Wolpe S.D., Masiarz F., Coit D., Cerami A.; "Resolution of the two components of macrophage inflammatory promacrophage inflammatory protein 1, and cloning and characterization of one of those components, macrophage inflammatory protein 1 beta."; J. Exp. Med. 168:2251-2259(1988).
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                                                                                                                                                                                                                                                                                                                                                   Chemotaxis; Inflammatory response; Signal.
Chemotaxis; Inflammatory response; Signal.
1 23 BY SIMILARITY.
24 92 SMALL INDUCIBLE CYTOKINE A4
34 58 BY SIMILARITY.
35 74 BY SIMILARITY.
92 AA; 10234 MW; 60B451EEEBC7103D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 81.1%; Score 394; DB 1; I larity 77.2%; Pred. No. 8.6e-37; Conservative 10; Mismatches 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-JAN-1990 (Rel. 13, Created)
01-FEB-1996 (Rel. 33, Last sequence update)
15-SEP-2003 (Rel. 42, Last annotation update)
Small inducible cytokine A4 precursor (CCL4)
protein 1-beta) (MIP-1-beta) (H400 protein) (3
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                                                                                                                                                                                                                                       EMBL; U06434; AAA96497.1; -.
HSSP; P13236; 1HUM.
InterPro; IPR000827; CC_chemkine_sml
InterPro; IPR001811; Chemokine_IL8.
Pfam; PF00048; IL8; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC
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                                                           (CHEMOKINE CC)
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Mammalia; Eutheria;
NCBI TaxID=10090;
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92 AA;
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SIGNAL
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                                                                                                          STRAIN=B10.S/J, and SJL/J; TISSUE=Spleen;
STRAIN=B10.S/J, and SJL/J; TISSUE=Spleen;
MEDLINE=99370037; PubMed=10438970;
Teuscher C., Butterfield R.J., Ma R.Z., Zachary J.F., Doerge R.W.,
Teuscher C., Butterfield R.J., Ma R.Z., Zachary J.F., Doerge R.W.,
Blankenhorn E.P.;
"Sequence polymorphisms in the chemokines Scyal (TCA-3), Scya2
(monocyte chemoattractant protein (MCP)-1), and Scyal2 (MCP-5) are
candidates for eae7, a locus controlling susceptibility to monophasic
remitting/nonrelapsing experimental allergic encephalomyelitis.";
J. Immunol. 163:2262-2266(1999).
-!- FUNCTION: MONOKINE WITH INFLAMMATORY AND CHEMOKINETIC PROPERTIES.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY (SMALL CYTOKINE
-:- SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY (SMALL CYTOKINE
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                                      Pragnell
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P10855; P14096;

01-JUL-1989 (Rel. 11, Created)

01-APR-1990 (Rel. 14, Last sequence update)

15-SEP-2003 (Rel. 42, Last annotation update)

Small inducible cytokine A3 precursor (CCL3) (Macriprotein 1-alpha) (MIP-1-alpha) (TY-5) (SIS-alpha)

chemotaxis protein) (L2G25B).
                                     Grove M.,
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Pred. No. 2.4e-36;
3; Mismatches 9;
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BY SIMILARITY.
BY SIMILARITY.
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rPro; IPROulc...

1; PF00048; IL8; 1.

1; SM00199; SCY; 1.

SITE; PS00472; SMALL CYTOKINES CC; 1

SITE; PS00472; SMALL CYTOKINES CC; 1

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EMBL; M35590; AAA39708.1; -.
EMBL; X62502; CAA44364.1; -.
EMBL; AF128218; AAF22559.1; -.
EMBL; AF128219; AAF22550.1; -.
PIR; C30552; C30552.
HSSP; P13236; 1HUM.
MGD; MGI:98261; Cc14.
InterPro; IPR000827; CC_chemkine_sm_InterPro; IPR001811; Chemokine_IL8.
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Similarity 76.1%;
70; Conservative
FROM N.A.
                      TRAIN=DBA/2J; TISSUE=Li
aubersies P., Lepretre
lumb M.A.;
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92 A5;
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InterPro; I
Pfam; PF000
SMART; SM00
PROSITE; PS
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SEQUENCE
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"Sequence polymorphisms in the chemokines Scyal (TCA-3), Scya2 (monocyte chemoattractant protein (MCP)-1), and Scyal2 (MCP-5) are candidates for eae7, a locus controlling susceptibility to monophasic remitting/nonrelapsing experimental allergic encephalomyelitis.";
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MEDLINE=88154745; PubMed=3279154;
Wolpe S.D., Davatelis G., Sherry B., Beutler B., Hesse D.G.,
Nguyen H.T., Moldawer L.L., Nathan C.F., Lowry S.F., Cerami A.;
Nguyen H.T., moldawer L.L., Nathan C.F., Lowry S.F., Cerami A.;
"Macrophages secrete a novel heparin-binding protein with
inflammatory and neutrophil chemokinetic properties.";
J. Exp. Med. 167:570-581(1988).
J. Exp. Med. 167:570-581(1988).
-!- FUNCTION: MONOKINE WITH INFLAMMATORY, PYROGENIC AND CHEMOKINETIC
-!- FUNCTION: MONOKINE WITH INFLAMMATORY, PYROGENIC AND CHEMOKINES.
BINDING TO A HIGH-AFFINITY RECEPTOR ACTIVATES CALCIUM RELEASE IN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN=DBA/2J;
MEDLINE=91016858; PubMed=2216738;
Grove M., Lowe S., Graham G., Pragnell I., Plumb M.;
"Sequence of the murine haemopoietic stem cell inhibitor/macrophage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Doerge R.W.,
                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE=88258380; PubMed=3290382;
MEDLINE=88258380; PubMed=3290382;
Davatelis G., Tekamp-Olson P., Wolpe S.D., Hermsen K., Luedke C.,
Gallegos C., Coit D., Merryweather J., Cerami A.;
Gallegos C., Coit D., Merryweather J., Cerami A.;
"Cloning and characterization of a cDNA for murine macrophage
"Cloning and characterization of a novel monokine with inflammatory and
inflammatory protein (MIP), a novel monokine with inflammatory and
chemokinetic properties.";
chemokinetic properties.";
                                               Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus
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Brown K.D., Zurawski S.M., Mosmann T.R., Zurawski G.;
Brown K.D., Zurawski S.M., Mosmann T.R., Zurawski G.;
"A family of small inducible proteins secreted by leukocytes are members of a new superfamily that includes leukocyte and fibroblast-derived inflammatory agents, growth factors, and indicators of various activation processes.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Genomic structure of murine macrophage inflammatory protein-1 and conservation of potential regulatory sequences with a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hermsen K., Luedke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
MEDLINE=91237116; PubMed=2033269;
Widmer U., Yang Z., van Deventer S., Manogue K.R., Sherry
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Б.
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86:1963-1967(1989)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in 1 alpha gene.";
18:5561-5561(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Davatelis G., Tekamp-Olson P., Wolpe Gallegos C., Coit D., Merryweather J. J. Exp. Med. 170:2189-2189(1989).
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STRAIN=BALB/CJ, DBA/2J, NOD/LTJ, S.
MEDLINE=99370037; PubMed=10438970;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
MEDLINE=89184547; PubMed=2784565;
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"cDNA sequences of two inducible
Proc. Natl. Acad. Sci. U.S.A. 86:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       homolog, LD78.";
J. Immunol, 146:4031-4040(1991)
                                                 Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10090;
OR MIP1A.
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                             musculus (Mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic Acids
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     OR SCYA3
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REVISIONS.
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is produced through a collaboration
                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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P16619; Q96168;
01-AUG-1990 (Rel. 15, Last sequence update)
01-AUG-1990 (Rel. 15, Last annotation update)
15-SEP-2003 (Rel. 42, Last annotation update)
Small inducible cytokine A3 like 1 precursor (Tonsillar lymphocyte LD78 beta protein) (G0/G1 switch regulatory protein 19-2) (G0S19-2 protein) (PAT 464.2).
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Catarrhini; Hominidae; Homo.
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EMBL: X12531; CAA31452.1; -..
EMBL: X12531; CAA37452.1; -..
EMBL: J04491; AAA40304.1; -..
EMBL: AF065939; AAC17506.1; -..
EMBL: AF065940; AAC17506.1; -..
EMBL: AF065941; AAC17500.1; -..
EMBL: AF065941; AAC17500.1; -..
EMBL: AF065941; AAC17500.1; -..
EMBL: AF065943; AAC17500.1; -..
EMBL: AF065941; AAC17500.1; -..
EMBL: 
                                                             SPLEEN, AND
BETA FAMILY
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No. 1.1e-28;
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Pred. No. 1.16
16; Mismatches
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SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: EXPRESSED IN LUNG,
SIMILARITY: BELONGS TO THE INTERCRINE 1
C-C) (CHEMOKINE CC).
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Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein) (PAT 464.2).
CCL3L1 OR SCYA3L1 OR G0S19-2.
Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        65.9%;
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56; Conserv
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TISSUE=Blood;
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Best Local
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SECULANCE C.

STRAIN=Wistar;

MEDLINE=96183056; PubMed=8607872;

MEDLINE=96183056; PubMed=8607872;

MEDLINE=96183056; PubMed=8607872;

Makagawa H., Shiota S., Takano K., Shibata F., Kato H.;

Nakagawa H., Shiota S., Takano K., Shibata F., Kato H.;

Makagawa H., Shiota S., Takano K., Shibata F., Kato H.;

Nakagawa H., Shiota S., Takano K., Shibata F., Kato H.;

Incytokine-induced neutrophil chemoattractant (CINC)-2 alpha, a novel of the member of rat GRO/CINCS, is a predominant chemokine produced by the member of rat GRO/CINCS Manual CASINGT INFLAMMATORY AND CHEMOKINETIC PROPERTIES.

Incommon Sequence of the member of the mem
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; Chordata; Craniata; Vertebrata; Buteleostomi;
; Rodentia; Sciurognathi; Muridae; Murinae; Rattus
                                                                                                                                                                                                                                                                                                                                                                        Gaps
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ein-1 alpha (MIP-1 alpha)
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E CRC64;
                                                                                                                        CYTOKINE A3
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TISSUE=Lung;

MEDLINE=95298037; PubMed=7779098;

Shi M.M., Godleski J.J., Paulauskis J.D.;

"Molecular cloning and posttranscriptional regulation o;

"Molecular cloning and posttranscriptional regulation o;

inflammatory protein-1 alpha in alveolar macrophages.";

biochem. Biophys. Res. Commun. 211:289-295(1995).
                                                                                                                                                                                                                                                                                                                Length
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STRAIN=Long Evans; TISSUE=Lung;
MEDLINE=95238980; PubMed=7722328;
Shanley T.P., Schmal H., Friedl H.P., Jones M.L.,
"Role of macrophage inflammatory protein-1 alpha acute lung injury in rats.";
J. Immunol. 154:4793-4802(1995).
                                                                                                                                                                                                                                                                                                          Score 320; DB 1; I pred. No. 1.3e-28; 0; Mismatches 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
01-OCT-1996 (Rel. 41, Last annotation update)
28-FEB-2003 (Rel. 41, Last annotation update)
Small inducible cytokine A3 precursor (CCL3) (Ms protein 1-alpha) (MIP-1-alpha).
CCL3 OR SCYA3 OR MIP1A.
Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebre Mammalia; Eutheria; Rodentia; Sciurognathi; Muri
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3Y LIPOPOLYSACCHARIDE (LPS).

BELONGS TO THE INTERCRINE BETA
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                                                                                                                        INDUCIBLE
                                                                                                                       SMALL INDUCIBLE
BY SIMILARITY.
BY SIMILARITY.
L -> P (IN REF.
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                           CC;
                               CYTOKINES
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SM00199; SCY; 1.
; PS00472; SMALL CYTO:
le; Chemotaxis; Signal
                                                                                                                                                                                                                                                                                                                55.84;
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SUBCELLULAR I
INDUCTION: B)
SIMILARITY: I
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F
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     SMART; SMO
PROSITE; P
Cytokine;
SIGNAL
CHAIN
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DISULFID
CONFLICT
SEQUENCE
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                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                   collaboration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-MAR-1989 (Rel. 10, Created)
01-MAR-1989 (Rel. 10, Last sequence update)
15-SEP-2003 (Rel. 42, Last annotation update)
Small inducible cytokine A3 precursor (CCL3) (Macrophage inflammatory protein 1-alpha) (MIP-1-alpha) (Tonsillar lymphocyte LD78 alpha protein) (G0/G1 switch regulatory protein 19-1) (G0S19-1 protein)
(SIS-beta) (PAT 464.1).
CCL3 OR SCYA3 OR G0S19-1 OR MIP1A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                             Signal; Heparin-binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             [2]
SEQUENCE FROM N.A.
MEDLINE=89140347; PubMed=2521882;
MEDLINE=89140347; PubMed=2521882;
Zipfel P.F., Balke J., Irving S.G., Kelly K., Siebenlist U.;
Zipfel P.F., Balke J., Irving S.G., Kelly K., Siebenlist U.;
Mitogenic activation of human T cells induces two closely related "Mitogenic activation of human T cells induces two closely related "Mitogenic activation of human gimilarities with a new family of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Euteleostomi
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MEDLINE=86223879; PubMed=3086300;
MEDLINE=86223879; PubMed=3086300;
Obaru K., Fukuda M., Maeda S., Shimada K.;
"A cDNA clone used to study mRNA inducible in human tonsillar lymphocytes by a tumor promoter.";
Jymphocytes by a tumor promoter.";
J. Biochem. 99:885-894(1986).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                               CYTOKINE A3
                                                                                                                                                                                                                                                                                                                                                            BY SIMILARITY.

BY SIMILARITY.

A -> T (IN REF. 2).

C -> W (IN REF. 2 AND 3).

14E861C647F9A2EB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 303.5; DB 1;
Pred. No. 8.3e-27;
7; Mismatches 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 91
                                                                                                                                                                                                                                                                                                                 response;
                                                                                                                                                                                                                                                                                                                                                 INDUCIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 PAVVFQTKRSKQVCADPSESWVQEYVYDLELN 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  60 PGVIFLTKRNRQICADPKETWVQEYITELELN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     92
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CC
                                                                                                                                                         EMBL; U22414; AAA80608.1; -.
EMBL; U06435; AAA96498.1; -.
PIR; I52322; I52322.
HSSP; P13236; IHUM.
InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_ILB.
Pfam; PF00048; ILB; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC; Cytokine; Chemotaxis; Inflammatory resignal.
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J. Immunol. 142:1582-1590(1989).
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10335 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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(CHEMOKINE
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                                                                                                                                                                                                                                                                                                                                                                                                                                    92 AA;
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Mammalia; Euther
NCBI TaxID=9606;
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4 4 6 6
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DISULFID
CONFLICT
CONFLICT
SEQUENCE
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE OF 27-40 AND 71-83, AND FUNCTION.

SEQUENCE OF 27-40 AND 71-83, AND FUNCTION.

MEDLINE=96106406; PubMed=8525373;

Cocchi F., DeVico A.L., Garzino-Demo A., Arya S.K., Gallo R.C.,

Lusso P.;

Lusu P. Location

Lusso P.;

Lusu P. Lumor PROMOTER); PHA = PHYTOHEMAGGLUTININ (T-CELL
                                                                                                                                                                                                                                                                                                                                                                                                     inflammatory protein-";
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                                              of
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                                                                                                                                                                                                                                                                                          SEQUENCE OF 24-92, AND MUTAGENESIS OF ASP-49.
MEDLINE=96127782; PubMed=8541527;
Hunter M.G., Bawden L., Brotherton D., Craig S., Cribbes S., Czaplewski L.G., Dexter T.M., Drummond A.H., Gearing A.H., Heyworth C.M., Lord B.I., Mccourt M., Varley P.G., Wood L.M., Edwards R.M., Lewis P.J.;
"BB-10010: an active variant of human macrophage inflammatory 1 alpha with improved pharmaceutical properties.";
Blood 86:4400-4408(1995).
                                                                                                                   SEQUENCE FROM N.A.
MEDLINE=90287155; PubMed=1694014;
Nakao M., Nomiyama H., Shimada K.;
"Structures of human genes coding for cytokine LD78 and their expression.";
                                              inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C:soluble fraction; TAS.
F:antiviral response protein activity;
F:chemokine activity; TAS.
F:signal transducer activity; TAS.
P:antibacterial humoral response (sensu
                                                                                                                                                                                                                            SEQUENCE OF 23-92 FROM N.A. Jang J.S., Kim B.E.; Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases
                                                   an
                                                encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   D.R.;
e gene
                                                                                                                                                                                                    Biol. 10:3646-3658(1990)
SEQUENCE FROM N.A.
MEDLINE=91103879; PubMed=2271120;
Blum S., Forsdyke R.E., Forsdyke Dalum S., Forsdyke Dalum S., Forsdyke Dalum S., Forsdyke Dalum S., Forsdyke Dalum Dony Cell Biol. 9:589-602(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CHEMOKINE CC). STRONG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; D00044; BAA00029.1; -.
EMBL; M23452; AAA36316.1; -.
EMBL; M25315; AAA57255.1; -.
EMBL; X03754; CAA27388.1; -.
EMBL; X04018; CAA27643.1; -.
EMBL; M23178; AAA35858.1; -.
EMBL; D90144; BAA14172.1; -.
EMBL; AF043339; AAC03539.1; -.
PIR; A35673; A30574.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B53; 22-JUL-99;
HGNC:10627; CC
82283; -.
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GO: 0003800;
GO: 0008009;
GO: 0004871;
GO: 0006961;
                                                                                                                                                                                  expression."
Mol. Cell. E
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PDB; 1B50;
PDB; 1B53;
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MIM, 18
GO; GO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
Hughes S.M., Bumstead N.;
"Mapping of the gene encoding the chicken homologue of the mammalian chemokine SCYA4.";
Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL/GenBank/DDBJ databases.
INFLAMMATORY AND CHEMOKINETIC PROPERTIES
                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=95369710; PubMed=7642115;
Petrenko O., Ischenko I., Enrietto P.J.;
"Isolation of a cDNA encoding a novel chicken chemokine homologous
mammalian macrophage inflammatory protein-1 beta.";
Gene 160:305-306(1995).
                                                                                                                                                                                                                                      response; Signal; 3D-structure
                                                                                                                                                                                                                                                                                                                                                                                                    .,
H
                                                                            protein signalin.
                                                                                                                                                                                                                                                                                                                                                                       92
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Q90826; Q910C9;
01-NOV-1997 (Rel. 35, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Small inducible cytokine A4 homolog precursor (Macrophage protein 1-beta homolog).
                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                               IMPROVED
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PHARMACEUTICAL PROPERTIES
517865D5D6776CA8 CRC64;
                                                                                                                                                                                                                                                                SMALL INDUCIBLE CYTOKINE BY SIMILARITY. BY SIMILARITY.
                                                                                                                      TAS
                                                                                                                                                                                                                                                                                                                                                                                                  19;
GO; GO: 0006874; P: calcium ion homeostasis; TAS.
GO; GO: 0006928; P: cell motility; TAS.
GO; GO: 0007267; P: cell-cell signaling; TAS.
GO; GO: 0006835; P: chemotaxis; TAS.
GO; GO: 0006887; P: exocytosis; TAS.
GO; GO: 0006955; P: immune response; TAS.
GO; GO: 0006955; P: inflammatory response; TAS.
GO; GO: 000711; P: regulation of cytoskeleton; TAS.
GO; GO: 0007165; P: signal transduction; TAS.
GO; GO: 0008166; P: viral replication; TAS.
InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_ILB.
FFam; PF00048; ILB; 1.
SMART; SM00199; SCY; 1.
                                                                                                                                                                                                                                                                                                                                                                        7;
                                                                                                                                                                                                                                                                                                                                                                       Score 300.5; DB 1
Pred. No. 1.8e-26;
); Mismatches 19
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; SM00199; SCY; 1.
; SM00199; SCY; 1.
TE; PS00472; SMALL CYTOKINES CC;
TE; PS00472; Inflammatory res
                                                                                                                                                                                                                                                                                                                                                                                                  19;
                                                                                                                                                                                                                                                                                                                                            10085 MW;
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hitted (JUL-1994) to the
FUNCTION: MONOKINE WITH
(BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                        61.8%;
larity 57.6%;
Conservative 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE OF 14-90 FROM N.A. Petrenko O., Enrietto P.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SCYA4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCL4 OR SCYA4.
Gallus gallus (Chic)
Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A. TISSUE=Bone marrow;
                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                               AA;
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9 4 8 4 9
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                                                                                                                                                                                                                                            Cytokine;
SIGNAL
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Best Local
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HUMAN
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Matches
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                                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See, http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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FROM
                       CYTOKIN
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Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordactyla; Ruminantia; Pecora; Bovoidea
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
                                                                                                                                                                                                                                           BY SIMILARITY.
SMALL INDUCIBLE CYTOKINE A4 HOMOLOG.
BY SIMILARITY.
BY SIMILARITY.
M -> L (IN REF. 1).
SOAF9679A267408F CRC64;
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TISSUE=Intestine;

Aust G., Thamm B., Rost A.K.;

Cloning of Bos taurus RANTES mRNA.";

Submitted (MAR-1999) to the EMBL/GenBank/DDBJ databases.

-!- FUNCTION: CHEMOATTRACTANT FOR BLOOD MONOCYTES, MEMORY TOTION: CHEMOATTRACTANT FOR BLOOD MONOCYTES, MEMORY TOTION: CHEMOATTRACTANT FOR BLOOD MONOCYTES, MEMORY TOTION BASOPHILS AND ACTIVATES EOSINOPHILS (BY SIMILARITY).

-!- SUBCELLULAR LOCATION: Secreted.

-!- SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY (SMAL
                        (SMALL
                                                                                                                                                                                                                                                                                                                          90
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                       FAMILY
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                                                                                                                                                                                                                                                                                                                                              23;
                                                                                                                                                                                                                                                                                                                         ore 264; DB 1;
ed. No. 1.8e-22;
Mismatches 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Small inducible cytokine A5 precursor (CCLS)
protein) (SIS-delta).
                          BETA
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                        INTERCRINE
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EMBL; AJ243034; CAB45103.1; -.
HSSP; P13236; 1HUM.
InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_ILB.
Pfam; PF00048; ILB; 1.
SMART; SM0199; SCY; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC; Cytokine; Chemotaxis; Signal.
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                                                                                                                                                                                                                         GG;
    similarity)
                                                                                                                                                                                                                                                                                                                           Score
Pred.
   SUBUNIT: Homodimer (By similar SUBCELLULAR LOCATION: Secrete SIMILARITY: BELONGS TO THE IN'C-C) (CHEMOKINE CC).
                                                                                                                                                                                                                                                                                                                                                20;
                                                                                                                                                                                                                                                                                           87
9987 MW;
                                                                                                                                                                                                                                                                                                                          th Similarity 51.1%; 45; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STANDARD;
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097919;
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PROSITE; P
Cytokine;
SIGNAL
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DISULFID
CONFLICT
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Best Local
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HCC-2, LEC, and
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SEQUENCE FROM N.A.
Zeng Q.P., Yang R.Y., Fu L.C.;
"The complete sequence of human beta-chemokine RANTES mRNA.";
"The complete sequence of human beta-chemokine RANTES mRNA.";
"The complete sequence of human beta-chemokine RANTES mRNA.";
                                                                                                        EMBL; AJ007043; CAAU...

R HSSP; P13501; 1RTN.

R InterPro; IPR000827; CC_chemkine_sml.

R InterPro; IPR001811; Chemokine_IL8.

R InterPro; IPR001811; Chemokine_IL8.

R SMART; SM00199; SCY; 1.

SMART; SM00199; SCY; 1.

PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.

PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.

PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.

POTENTIAL.

SMALL INDUCIBLE CYTOKINE A5.

TOTAL INDUCIBLE CYTOKINE A5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 246.5; DB 1; Length
Pred. No. 1.6e-20;
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Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=99228475; PubMed=10213461;
Nomiyama H., Fukuda S., Iio M., Tanase S., Miura R., Y
"Organization of the chemokine gene cluster on human of 17q11.2 containing the genes for CC chemokine MPIF-1, RANTES.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dyer B.J., Jorgensen J.,
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MEDLINE=88285659; PubMed=2456327;
Schall T.J., Jongstra J., Dyer B.J.,
Davis M.M., Krensky A.M.;
"A human T cell-specific molecule is
J. Immunol. 141:1018-1025(1988).
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
             MEDLINE=22388257; PubMed=12477932; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D., Altischul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K., Hockins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J., Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez M., Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E., Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E., Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
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Hoover D.M., Shaw J., Gryczynski Z., Proudfoot A.E.I., Wells T.N.C.,
Lubkowski J.;
Lubko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AND
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PubMed=8525373;
A.L., Garzino-Demo A., Arya S.K., Gallo R.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SYNTHESIS, AND X-RAY CRYSTALLOGRAPHY (1.6 ANGSTROMS).
MEDLINE=99111238; PubMed=9889151;
Wilken J., Hoover D., Thompson D.A., Barlow P.N., McSparron H. Picard L., Wlodawer A., Lubkowski J., Kent S.B.;
"Total chemical synthesis and high-resolution crystal structur the potent anti-HIV protein AOP-RANTES.";
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MEDLINE=95352612; PubMed=7542919;
Chung C.-W., Cooke R.M., Proudfoot A.E.I., Wells T.N.C.
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[8]
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"Proton NMR assignments and solution conformation of RAI chemokine of the C-C type.";
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MEDLINE=96106406; PubMed=8525373;
Cocchi F., DeVico A.L., Garzino-Demo A., Arya S.K.
Lusso P.;
"Identification of RANTES, MIP-1 alpha, and MIP-1
HIV-suppressive factors produced by CD8+ T cells."
Science 270:1811-1815(1995).
PubMed=12477932;
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cell; Signal; Inflammatory response
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GO; GO:0003800; F:antiviral response protein activity; TAS

GO; GO:0008009; F:chemokine activity; TAS.

GO; GO:0006928; P:cell adhesion; TAS.

GO; GO:0006928; P:cell adhesion; TAS.

GO; GO:0006928; P:cell-cell signaling; TAS.

GO; GO:0006968; P:cell-cell signaling; TAS.

GO; GO:0006968; P:cell-cell signaling; TAS.

GO; GO:0006969; P:cell-cell signaling; TAS.

GO; GO:0006969; P:cell-cell signaling; TAS.

GO; GO:0006979; P:response to oxidative stress; TAS.

GO; GO:0006979; P:response to oxidative stress; TAS.

GO; GO:0006979; P:response to viruses; TAS.

GO; GO:0007165; P:signal transduction; TAS.

InterPro; IPR001811; Chemokine sml.

InterPro; IPR001811; Chemokine ILB.

SMART; SM00199; SCY; 1.

PROSITE; PS00472; SMALL CYTOKINES CC; 1.

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7: T-cell; 6
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EMBL; AF043341; AAC03541.1;
EMBL; AF088219; AAC63331.1;
EMBL; AF266753; AAF73070.1;
EMBL; BC008600; AAH08600.1;
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; 1HRJ; 14-OCT-96.
; 1RTN; 03-JUN-95.
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; 1B3A; 23-APR-99.
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ew; HGNC:10632; CCL5.
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SEQUENCE FROM N.A.

STRAIN=C57BL/6J; TISSUE=Pancreas;

PubMed=12466851;

A Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kondo S.,

Nikaido I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H.,

A Yagi K., Tomaru Y., Hasegawa Y., Nogami A., Schonbach C., Gojobori T.,

Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,

Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,

A Schriml L.M., Kanapin A., Matsuda H., Batalov S., Beisel K.W.,

A Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,

Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,

A Gasterland T., Gariboldi M., Gissi C., Godzik A., Gough J.,

Grimmond S., Gustincich S., Hirokawa N., Jackson I.J., Jarvis E.D.,

A Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,
                                                                                                                                                                                                                                                                                                    cDNA from renal tubular epithelium
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SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
STRAIN=NIH Swiss;
MEDLINE=94132613; PubMed=7507961;
Danoff T.M., Lalley P.A., Chang Y.S., Heeger P.S., Neilson E.G.;
Danoff T.M., calley P.A., chang Y.S., Heeger P.S., Neilson E.G.;
"Cloning, genomic organization, and chromosomal localization of the
"Cloning, genomic organization, and chromosomal localization of the
scyas gene encoding the murine chemokine RANTES.";
J. Immunol. 152:1182-1189(1994).
                                          specific RANTES
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STRAIN=B10.S/J, BALB/CJ, DBA/2J, NOD/LTJ, and SJL/J; TISSUE=Spleen; STRAIN=B10.S/J, BALB/CJ, DBA/2J, NOD/LTJ, and SJL/J; TISSUE=Spleen; MEDLINE=99370037; PubMed=10438970;
Teuscher C., Butterfield R.J., Ma R.Z., Zachary J.F., Doerge R.W., Blankenhorn E.P.;
"Sequence polymorphisms in the chemokines Scyal (TCA-3), Scya2 (monocyte chemoattractant protein (MCP)-1), and Scya12 (MCP-5) are candidates for eae7, a locus controlling susceptibility to monophase remitting/nonrelapsing experimental allergic encephalomyelitis.";
J. Immunol. 163:2262-2266(1999).
                                                                                                                              Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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MEDLINE=92289805; PubMed=1376260;
Schall T.J., Simpson N.J., Mak J.Y.;
Schall T.J., Simpson N.J., Mak J.Y.;
"Molecular cloning and expression of the murine RANTES cytokine:
structural and functional conservation between mouse and man.";
Eur. J. Immunol. 22:1477-1481(1992).
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STRAIN=BALB/c;
MEDLINE=94217689; PubMed=7513046;
Shin H.S., Drysdale B.E., Shin M.L., Noble P.W., Fisher Shin H.S., Drysdale B.E., Shin M.L., Noble P.W., Fisher Paznekas W.A.;
"Definition of a lipopolysaccharide-responsive element if lanking regions of MuRantes and crg-2.";
flanking regions of MuRantes and crg-2.";
Mol. Cell. Biol. 14:2914-2925(1994).
                                            (T-cell
                                                                                                                                                                                                                                                               O'Farrell
01-JUN-1994 (Rel. 29, Last sequence update)
15-SEP-2003 (Rel. 42, Last annotation update)
Small inducible cytokine A5 precursor (CCLS)
protein) (SIS-delta) (MuRantes).
CCLS OR SCYAS.
                                                                                                                                                                                                                                       990; PubMed=1375672;
1f G., Meyers C., Sun M.J.,
Neilson E.G.;
                                                                                                                                                                                                                                                                 Heeger P., Wolf G., Meyers C., Sun Krensky A.M., Neilson E.G.; "Isolation and characterization of encoding murine Rantes."; Kidney Int. 41:220-225(1992).
                                                                                                        Mus musculus (Mouse)
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10090;
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"Organization of
for ClO, MRP-2
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STRAIN=129/Sv;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pubmed=124//932;
A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
A Strausberg R.L., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
Biatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
Brownstein M.J., Widdin T.B., Toshiyuki S., Carninci P., Prange C.,
Brownstein M.J., Widdin T.B., Toshiyuki S., Carninci P., Prange C.,
Brownstein M.J., Widdin T.B., Toshiyuki S., Carninci P., Prange C.,
Brownstein M.J., McKernan K.J., Malek J.A., Gunaraten P.H.,
Bosak S.A., McEwan P.J., McKernan K.J., Marek J.A., Gibbs R.A.,
A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
A Nilalon D.K., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
Butterfield X.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
Butterfield X.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
Generation and initial analysis of more than 15,000 full-length human
Konagaya A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,

Maglott D.R., Maltais L., Marchionni L., McKenzie L., Miki H.,

Nagashima T., Numata K., Okido T., Pavan W.J., Pertea G., Pesole G.,

Ravasi T., Reed J.C., Reed D.J., Reid J., Ring B.Z., Ringwald M.,

Sandelin A., Schneider C., Semple C.A., Setou M., Shimada K.,

Sultana R., Takenaka Y., Taylor M.S., Teasdale R.D., Tomita M.,

Verardo R., Wagner L., Wahlestedt C., Wang Y., Watanabe Y., Wells C.,

Wilming L.G., Wynshaw-Boris A., Yanagisawa M., Yang L.,

Wilming L.G., Wynshaw-Boris A., Yanagisawa M., Yang L.,

Minaki T., Waki K., Konno H., Nakawura M., Sakazume N., Sato K.,

A Hirozane-Kishikawa T., Konno H., Nakawura M., Sakazume N., Sato K.,

A Hara A., Hashizume W., Imotani K., Ishii Y., Itoh M., Kagawa I.,

Myazaki A., Sakai K., Sasaki D., Shibata K., Shinagawa A.,

A yasunishi A., Yoshino M., Waterston R., Lander E.S., Rogers J.,

Birney E., Hayashizaki Y.,

"Analysis of the mouse transcriptome based on functional annotation of

60,770 full-length CDNAS.";

Nature 420:563-573(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).

-! FUNCTION: CHEMOATTRACTANT FOR BLOOD MONOCYTES, MEMORY T HELPER
CELLS AND EOSINOPHILS. CAUSES THE RELEASE OF HISTAMINE FROM
BASOPHILS AND ACTIVATES EOSINOPHILS.

-! SUBCELLULAR LOCATION: Secreted.

-! SUBCELLULAR LOCATION: T-CELL AND MACROPHAGE SPECIFIC.

-! TISSUE SPECIFICITY: T-CELL AND MACROPHAGE SPECIFIC.
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TISSUE=Mammary gland;
PubMed=12477932;
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; BC033508; I48875; A4 P13501;

EMBL;

EMBL;

PIR; I HSSP;

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STANDARD; PRT; 91 AA.

0912L1;

28-FEB-2003 (Rel. 41, Last annotation update)

28-FEB-2003 (Rel. 41, Last annotation update)

Small inducible cytokine A5 precursor (CCL5) (T-cell specific RANTES)

protein).

CCL5 OR SCYAS.

Sigmodon hispidus (Hispid cotton rat).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Sigmodontinae;

Sigmodon.

NCBI_TaxID=42415;
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SEQUENCE FROM N.A.
Stance J.C., Pletneva L.M., Prince G.A.;
Sigmodon hispidus cytokines, chemokines and interferons.";
Submitted (SEP-2001) to the EMBL/GenBank/DDBJ databases.
Submitted (SEP-2001) to the EMBL/GenBank/DDBJ databases.
CELLS AND EOSINOPHILS. CAUSES THE RELEASE OF HISTAMINE BASOPHILS AND ACTIVATES EOSINOPHILS (BY SIMILARITY).

SASOPHILS AND ACTIVATES EOSINOPHILS (SY SIMILARITY).

SUBCELLULAR LOCATION: Secreted.
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embl; AF421391; AAL16932.1; ...
EMBL; AF421391; AAL16932.1; ...
InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_IL8.

SMART; SM00199; SCY; 1.

SMART; SM00199; SCY; 1.

R PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.

R PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.

Cytokine; Chemotaxis; T-cell; Signal; Inflammatory reserriche; Chemotaxis; T-cell; Signal; Inflammatory reserriches.

Z4 91 SMALL INDUCIBLE CYTOKINE
BY SIMILARITY.
                                                                                                                                                                                                                              CYTOKINE
                                                                                                                                                                                                                                                                                                                                                      -> E (IN REF. 1).
5DFD66F4684FE1C8 CRC64;
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48.9%; Pred. No. 4.4e-20;
ive 20; Mismatches 26;
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F-cell; Signal; Inflamma
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                              chemkine sml
MGD; MGI:98262; Cc15.
InterPro; IPR000827; CC_chemkine_InterPro; IPR001811; Chemokine_II_Pfam; PF00048; IL8; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL_CYTOKINESCYTOKINESCYTOKINE; Cytokine; Chemotaxis; T-cell; Signation of the context of 
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                                                                                                                     MKLCVTVLSLLMLVAAFCSPALSAPMGSDPPTACCFSYTARKLPRNFVVDYYETSSLCSQ
                                                                                                                                              1 MKISAAVLTVVLMAASLCAPASASPNGSD-TIPCCFAYLSAVLPRAHVKEYFYTSSKCSN
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                                                                                                                                                                                                                                                                                                                                                                                                            (T-cell specific RANTES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUE=Lung;
Asano K., Nakamura M., Oguma T., Fukunaga K., Ishizaka A.,
Yamaguchi K., Kanazawa M.;
Yamaguchi K., Kanazawa M.;
Submitted (APR-1997) to the EMBL/GenBank/DDBJ databases.
-:- FUNCTION: CHEMOATTRACTANT FOR BLOOD MONOCYTES, MEMORY T HELPER
CELLS AND EOSINOPHILS. CAUSES THE RELEASE OF HISTAMINE FROM
BASOPHILS AND ACTIVATES EOSINOPHILS (BY SIMILARITY).
-:- SUBCELLULAR LOCATION: Secreted.
-:- SUBCELLULAR LOCATION: Secreted.
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EMBL; U77037; AAC53293.1;

R HSSP; P13501; 1RTN.

R InterPro; IPR000827; CC_chemkine_sml.

R InterPro; IPR010811; Chemokine_IL8.

DR Pfam; PF00048; 1L8; 1.

DR SMART; SM00199; SCY; 1.

PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.

PROSITE; PS00472; SMALL_CYTOKINE AS.

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RY31B751237DB9 CRC64;

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STRAIN=Dunkin-Hartley;
Campbell E.M., Proudfoot A.E.I., Yoshimura T., Allet B.,
Wells T.N.C., White A.M., Westwick J., Watson M.L.;
Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cavia porcellus (Guinea pig).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
Mammalia; Eutheria; Rodentia; Hystricognathi; Cavildae;
NCBI_TaxID=10141;
               DODGEAEABE4242FF CRC64;
                                               DB 1;
                                                                                   25;
                                                 49.1%; Score 238.5; DB 1 49.5%; Pred. No. 1.2e-19;
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P97272; 009076;
01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Small inducible cytokine A5 precursor (CCL5)
protein) (SIS-delta).
                                                                                                                                                                                             61 PAVVFQTKRSKQVCADPSESWVQEYVYDLEL 91
                                                                                  Mismatches
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10082 MW;
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SEQUENCE FROM N.A.
STRAIN=Long Evans; TISSUE=Lung;
Jones M.L., Shanley T.P., Ward P.A.;
Jones M.L., Shanley T.P., Ward P.A.;
Submitted (FEB-1994) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: CHEMOATTRACTANT FOR BLOOD MONOCYTES, MEMORY T HELPER
-!- FUNCTION: CHEMOATTRACTANT FOR BLOOD MONOCYTES, MEMORY T HELPER
-!- SUBCELLULAR INTERCRINE BETA FAMILY (SMALL CYTOKIN:
-!- SIMILARITY: BELONGS TO THE INTERCRINE BETA FAMILY (CHEMOKINE CC).
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STANDARD; PRT; 92 AA.

1 P50231;

1 01-0CT-1996 (Rel. 34, Created)

1 28-FEB-2003 (Rel. 41, Last annotation update)

1 28-FEB-2003 (Rel. 41, Last annotation update)

1 28-FEB-2003 (Rel. 41, Last annotation update)

2 Small inducible cytokine A5 precursor (CCL5) (T-cell specific RANTES Expotein) (SIS-delta).

8 Fattus norvegicus (Rat).

9 CCL5 OR SCYAS.

9 Rattus norvegicus (Rat).

9 Rattus norvegicus (Rat).

9 Rattus norvegicus (Rat).

9 Rattus norvegicus (Rat).

9 AA.
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d. No. 3.3e-19;
Mismatches 28;
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60 LAUVFVTRKNRQVCANPEKKWVREYINSLEMS
    Pred
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Q8hyq2 macaca mula
Q91zl0 sigmodon hi
Q8sqa6 bos taurus
Q91z65 sigmodon hi
Q8hyq3 macaca mula
Q14745 homo sapien
Q9pwa6 gallus gall
Q8hyn3 macaca mula
Q8hyn3 macaca mula
Q8hyq1 macaca mula
Q8mkd0 equus cabal
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Q91ZE5
Q91Z65
Q8HYQ3
Q14745
Q9PWA6
Q8HYN3
Q8HYN4
Q918E0
Q8HYQ1
Q8HYQ1
Q8HYQ1
Q8HYS0
Q8HYS0
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"Variation in gene copy number of the human chemokines macrophage inflammatory protein-
Inflammatory protein-la/CCL3 and macrophage inflammatory protein-
Ib/CCL4.";
Submitted (FEB-2002) to the EMBL/GenBank/DDBJ databases.

EMBL; AY079147; AAL87008.1; -.

EMBL; AY079147; AAL87008.1; -.

InterPro; IPR001811; Chemokine_IL8.

InterPro; IPR001811; Chemokine_IL8.

R Ffam; PF00048; IL8; 1.

R SMART; SM00199; SCY; 1.

R PROSITE; PS00472; SMALL CYTOKINES CC; 1.

R PROSITE; PS00472; SMALL CYTOKINES CC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                         Homo sapiens (Human).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Pred. No. 1.8e-49;
1; Mismatches 2; Indels
                                          OBNHW4;
01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
Macrophage inflammatory protein-1b2.
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Best Local Similarity
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; Sigmodontinae;
                                                                                                                                     eostomi;
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                                                                                                                                                                                                                     SEQUENCE FROM N.A.

Basu S., Schaefer T.M., Ghosh M., Fuller C.L., Reinhart T.A.;

Basu S., Schaefer T.M., Ghosh M., Fuller C.L., Reinhart T.A.;

"Comprehensive cloning and sequencing reveals evolutionary
conservation among all groups of rhesus macaque chemokines.";

Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases.

EMBL; AF449267; AAN76071.1; -.

EMBL; AF449267; AAN76071.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

Blanco J.C., Pletneva L.M., Prince G.A.;

Blanco J.C., Pletneva L.M., Prince G.A.;

Sigmodon hispidus cytokines, chemokines and interferons.";

Submitted (SEP-2001) to the EMBL/GenBank/DDBJ databases.

EMBL; AF421392; AAL16933.1; -.

HSSP; Q9Y258; 1G2S.

InterPro; IPR000827; CC_chemkine_sml.

InterPro; IPR001811; Chemokine_ILB.

R Pfam; PF00048; ILB; 1.

SMART; SM00199; SCY; 1.

R PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.

R PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.

SROUENCE 92 AA; 10195 MW; A34FDE21E6FA9C2E CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92;
                                                                                                                                    Vertebrata; Eutele; Cercopithecidae,
                                                                                                                                                                                                                                                                                                                                                92
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         update)
on update)
                                                                         Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                Score 459; DB 6; I pred. No. 2.3e-48; 3; Mismatches 3;
                                                                      update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            h Similarity 71.7%; Score 376; DB 11; Similarity 71.7%; Pred. No. 3.1e-38; 66; Conservative 13; Mismatches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     O91ZLO;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence up
01-MAR-2003 (TrEMBLrel. 23, Last annotation
Macrophage inflammatory protein 1 beta.
MIP-1BETA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92
                                                  01-MAR-2003 (TrEMBLrel. 23, Created)
01-MAR-2003 (TrEMBLrel. 23, Last sequence up
01-MAR-2003 (TrEMBLrel. 23, Last annotation
Chemokine CCL4/MIP-IBETA.
Macaca mulatta (Rhesus macaque).
Eukaryota; Metazoa; Chordata; Craniata; Vert
Mammalia; Eutheria; Primates; Catarrhini; Ce
Cercopithecinae; Macaca.
NCBI TaxID=9544;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94.4%;
93.5%;
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Euteleostomi;
; Sigmodontinae;
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01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-JUN-2002 (TrEMBLrel. 22, Last annotation update)
Macrophage inflammatory protein-1 alpha.
Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
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SEQUENCE FROM N.A.

Blanco J.C., Pletneva L.M., Prince G.A.;

"Sigmodon hispidus cytokines, chemokines and interferons.";

Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases.

EMBL; AY059407; AAL26704.1; -.

HSSP; Q9Y258; 1G28.

InterPro; IPR000827; CC_chemkine_sml.

InterPro; IPR001811; Chemokine_IL8.

InterPro; IPR001811; Chemokine_IL8.

SMART; SM00199; SCY; 1.

PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.

SEQUENCE 92 AA; 10334 MW; CF9AAB3D94DCAF79 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22; Indels
                                                                                                                                                                                                                                                                                                                                  "Role of chemokines in RSV infection.";
Submitted (JAN-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AY077840; AAL78060.1; -.
InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_IL8.
InterPro; IPR001811; Chemokine_IL8.
SMART; SM00199; SCY; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL CYTOKINES CC; 1.
PROSITE; PS00472; SMALL CYTOKINES CC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sigmodon hispidus (Hispid cotton rat).
Sigmodon hispidus (Hispid cotton rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
Macrophage inflammatory protein-1 alpha.
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Pred. No. 2.2e-31;
4; Mismatches 22,
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61 PAVVPQTKRSKQVCADPSESWVQEYVYDLELN
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llarity 60.9%;
Conservative 1
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NCBI_TaxID=9913;
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56; Conserv
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NCBI_TaxID=42415;
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Yama H., Iio M., Miura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Basu S., Schaefer T.M., Ghosh M., Fuller C.L., Reinhart T.A "Comprehensive cloning and sequencing reveals evolutionary conservation among all groups of rhesus macaque chemokines. Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases. EMBL; AF449266; AAN76070.1; -. SEQUENCE 92 AA; 10119 MW; 021CAA371143D12A CRC64;
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Q14745
Q1475;
Q1475;
Q1-NOV-1996 (TrEMBLrel. 01, Created)
Q1-NOV-1996 (TrEMBLrel. 01, Last sequence update)
Q1-NOV-1996 (TrEMBLrel. 20, Last annotation update)
LD78 alpha beta precursor (Fragment).
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
NCBI_TaxID=9606;
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Pred. No. 1.1e-26;
2; Mismatches 21;
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Miyakawa T.;
Submitted (AUG-1995) to the EMBL/GenBank/DDBJ da EMBL; D63785; BAA09855.1; -.
HSSP; P13236; 1HUM.
InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_IL8.
Pfam; PF00048; IL8; 1.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.
Pred. No. 2.8e-28;
; Mismatches 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92
                                                                                                                                                                                                                     92
                                                                                                                                                                                                                                                                               4
                                                                                                                                                                                                                                                                                                                                                                                                                                                             OBHYO3;

OBHYO3;

O1-MAR-2003 (TrEMBLrel. 23, Last sequence up

O1-MAR-2003 (TrEMBLrel. 23, Last annotation

O1-MAR-2003 (TrEMBLrel. 23, Last annotation

Chemokine CCL3/MIP-1ALPHA.

Macaca mulatta (Rhesus macaque).

Eukaryota; Metazoa; Chordata; Craniata; Vert

Mammalia; Eutheria; Primates; Catarrhini; Ce

Cercopithecinae; Macaca.
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TISSUE=Brain;
Ishizuka K., Igata-Yi R., Naruse K., Naka:
Katsuragi S., Kin Y., Ohmoto Y., Nomiyama
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60 PGVIFLIKRGRQVCADPSKEWVQKYVSDLELS
                                                                                                                                                                                                                   PAVVFOTKRSKOVCADPSESWVQEYVYDLELN
                                                                                                                                                                                                                                                   60 PGIIFLTKRNRHVCADPKETWVQEIITDLELN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22;
                            19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  th Similarity 52.2%; 48; Conservative 2
     Similarity 54.3%; 0; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A. Basu S., Schaefer "Comprehensive clo
                               50;
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                                                                                                                                            67
                                                                                                                                                                       9
                                                                                                                                          3 VSVAALAVLLIAICYQTSAAPVGSDPPTSCCFTYISRQLPFSFVADYYETNSQCPHAGVV
                                                                                                                                                                                                                                                                                                                                                                                                   en/.
Chordata; Craniata; Vertebrata; Euteleostomi;
Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.

Hughes S.M., Bumstead N.;

Hughes S.M., Bumstead N.;

"Mapping of the gene encoding the chicken homologue of the mammalian chemokine SCYA4.";

Submitted (APR-1999) to the EMBL/GenBank/DDBJ databases.

EMBL; AF146730; AAD48772.1; -.

HSSP; P13236; 1HUM.

InterPro; IPR000827; CC_chemkine_sml.

InterPro; IPR001811; Chemokine_IL8.

Pfam; PF00048; IL8; 1.

SMART; SM00199; SCY; 1.

PROSITE; PS00472; SMALL CYTOKINES CC; 1.

PROSITE; PS00472; SMALL CYTOKINES SOAF9679A26751CB CRC64;
                                                                                                                   Gaps
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
Macrophage inflammatory protein 1 beta (Fragment).
Macaca mulatta (Rhesus macaque).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
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                                                                                                                   ۲,
                                                                                          Length 80;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                54.3%; Score 264; DB 13; Length 5.larity 51.1%; Pred. No. 1.5e-24; Conservative 20; Mismatches 23; Indels
                                                                                                                    Indels
                                                                8B509EB15648E971 CRC64;
                                                                                                                                                                                                                                                                                                                          (Tremblrel. 13, Created)
(Tremblrel. 13, Last sequence update)
(Tremblrel. 20, Last annotation update)
                                                                                          DB 4;
                                                                                                                   14;
                                                                                          Score 271.5; DB 4
Pred. No. 1.6e-25;
                                    LD78 ALPHA BETA.
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                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            63 FITRKGREVCANPONDWVQDYMNKMELN
                          POTENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
                                                                                                                                                                                                    68 KRSKQVCADPSESWVQEYV 86
                                                                                                                                                                                                                          61 KRGRQVCADPSEEWVQKYV 79
                                                                                                                    17;
                                                                                                                                                                                                                                                                                                                                                                                          Gallus gallus (Chicken).
Eukaryota, Metazoa, Chordata, C
Archosauria, Aves, Neognathae,
            1
16
>80
80
8857 MW;
                                                                                            55.9%;
59.5%;
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                                                                                                                       Conservative
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SEQUENCE FROM N.A
                                                                 80 AA;
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01-MAR-2003
01-MAR-2003
01-MAR-2003
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01-MAY-2000
01-MAR-2002
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Best Local S
Matches 45
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SEQUENCE
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Signal.
NON TER
SIGNAL
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Q9PWA6;
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Gallus
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Matches
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Q9PWA6
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Q8HYN3
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Conservative
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                                                                                                                          21
89
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                                                                                                                                             89 AA;
                                                                                                                                   22
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01-MAR-2003
01-MAR-2003
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Best Local S
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                       SEQUENCE FROM N.A.
Hofmann-Lehmann R., Williams A.L., Swenerton R.K., Li P.-L.,
Hofmann-Lehmann R., Williams A.L., McClure H.M., Ruprecht R.M.;
Rasmussen R.A., Chenine A.-L., McClure H.M., Ruprecht R.M.;
"Quantification of Simian Cytokine and Beta-Chemokine mRNAs using
Real-Time Reverse-Transcriptase Polymerase Chain Reaction: Variations
in Expression during Chronic Primate Lentivirus Infection.";
Submitted (DEC-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AF457196; AAN76986.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                            tions
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  86
                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
Hofmann-Lehmann R., Williams A.L., Swenerton R.K., Li P.-L.,
Rasmussen R.A., Chenine A.-L., McClure H.M., Ruprecht R.M.;
"Quantification of Simian Cytokine and Beta-Chemokine mRNAs using
Real-Time Reverse-Transcriptase Polymerase Chain Reaction: Variati
in Expression during Chronic Primate Lentivirus Infection.";
Submitted (DEC-2001) to the EMBL/GenBank/DDBJ databases.
NON_TER
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Chordata, Craniata, Vertebrata, Euteleostomi,
Jeognathae, Galliformes, Phasianidae, Phasiani
                                                                                                                                                                                                                                                                                                                                  Euteleostomi
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                                                                                                                                                                                            83
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                                                                                                                                                                                       CFSYTARKLPRNFVVDYYETSSLCSQPAVVFQTKRSKQVCADPSESWVQ
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
Macrophage inflammatory protein 1 alpha (Fragment).
Macaca mulatta (Rhesus macaque).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutelec Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
NCBI_TaxID=9544;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         56;
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                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           update)
                                                                                                                                                Score 252; DB 6; 1
Pred. No. 2.2e-23;
; Mismatches 2;
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9
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Last sequence up
Last annotation
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No. 6
                                                                                                                                                                                                                                                              PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Cri
Archosauria; Aves; Neognathae; Gallus.
NCBI_TaxID=0^2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10;
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                                                                                                                                                similarity 93.9%;
46; Conservative
                                                                                                                              5663 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6442 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q918E0,
Q918E0,
01-OCT-2000 (TrEMBLrel. 15
01-OCT-2001 (TrEMBLrel. 15
01-DEC-2001 (TrEMBLrel. 15
Chemokine K203 precursor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                              PRELIMINARY;
 ercopithecinae, Macaca
CBI TaxID=9544;
                                                                                                                      4
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41; Conserv
                                                                                                                              49 AA;
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                                                                                    Submitted (DEC-20
EMBL; AF457196; ANON TER 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
Takafuji V.A., Sharova L.V., Crisman M.V., Howard R.D.;
Takafuji V.A., Sharova L.V., Crisman M.V., Howard R.D.;
"Equus caballus RANTES mRNA.";
Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AF506970; AAM34212.1; -.
InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_IL8.
InterPro; IPR001811; Chemokine_IL8.
SMART; SM00199; SCY; 1.
PROSITE; PS00472; SMALL_CYTOKINES_CC; 1.
SEQUENCE 91 AA; 10159 MW; AEBZ53E8CD4ED7FD CRC64;
                                                                                                                                                                                                                                                                                                                                            DB 13; Length
                                                                                                                                                                                                                                                                                                                                                                                  Indels
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                                                                                                                                                                                                                                                      POTENTIAL.
CHEMOKINE K203.
6FA2EA7A4950CA75 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota, Metazoa, Chordata, Craniata, Vertebrata,
Mammalia, Eutheria, Perissodactyla, Equidae, Equus.
NCBI_TaxID=9796;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      update)
                                                                                                                                                                                                                                                                                                                         Score 245.5; DB 13;
Pred. No. 2.6e-22;
                                                                       Weining K.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q8MKD0;
01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation updat
Small inducible cytokine A5 RANTES.
Equus caballus (Horse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 6;
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23, Last sequence update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PAVVFOTKRSKQVCADPSESWVQEYVYDLELN 92
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SEQUENCE FROM N.A.

MEDLINE=20170941; PubMed=10704244;
Sick C., Schneider K., Staeheli P., Wein
"Novel chicken CXC and CC chemokines.";
Cytokine 12:181-186(2000).
EMBL; Y18692; CAB70956.1; -.
HSSP; P13236; 1HUM.
InterPro; IPR001811; Chemokine_IL8.
Pfam; PF00048; IL8; 1.
SMART; SM00199; SCY; 1.
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MEDLINE=21655115; PubMed=11797102;

Hughes S., Haynes A., O'Regan M., Bumstead N.;

"Identification, mapping, and phylogenetic analysis of three novel

T chicken CC chemokines.";

I mmunogenetics 53:674-683(2001).

R EMBL; AY037859; AAK84432.1; -.

InterPro; IPR000827; CC chemkine sml.

R InterPro; IPR001811; Chemokine_IIB.

R Pfam; PF00048; ILB; 1.

R SMART; SM00199; SCY; 1.

R PROSITE; PS00472; SMALL CYTOKINES CC; 1.

R PROSITE; PS00472; SMALL CYTOKINES CC; 1.
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                                                                                                              Basu S., Schaefer T.M., Ghosh M., Fuller C.L., Reinhart T.A.; "Comprehensive cloning and sequencing reveals evolutionary conservation among all groups of rhesus macaque chemokines."; Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases.

EMBL; AF449268; AAN76072.1; -. SEQUENCE 91 AA; 10019 MW; BCBD69ED0573803B CRC64;
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01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
Chemokine CCL5/RANTES.
Macaca mulatta (Rhesus macaque).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutelec
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
Cercopithecinae; Macaca.
NCBI_TaxID=9544;
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MW; 744A64BB229194EF CRC64;
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Last annotation update)
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Pred. No. 1.1e-21;
3; Mismatches 26;
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29;
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Q8QG57;
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Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; CreArchosauria; Aves; Neognathae; Gallus.
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larity 45.7%;
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Enomoto A., Kano R., Hasegawa A.;

"molecular cloning of canine RANTES gene.";

Submitted (DEC-2002) to the EMBL/GenBank/DDBJ databases.

EMBL; AB098562; BAC53725.1; -.

SEQUENCE 91 AA; 10179 MW; 01D79538CB8148E7 CRC64;
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01D79538CB8148E7 CRC64;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Carnivora; Fissipedia; Canidae;
NCBI_TaxID=9615;
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                                                                                                                                                                                                                                                                 DB 6;
                                                                                                                                                                                                                                                                  Score 225.5; DB 6;
Pred. No. 7.4e-20;
1; Mismatches 29;
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US-08-390-740B-7
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US-09-081-230-3
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US-09-113-705-4
US-09-113-705-7
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PCT-US94-08207A-28
Sequence 28, Application PC/TUS9408207A
Sequence 28, Application PC/TUS9408207A
GENERAL INFORMATION:
APPLICANT: The Upjohn Company
APPLICANT: FOR U.S. PURPOSES ONLY: Hoogewerf, Arlene J. a
ITILE OF INVENTION: USE OF HEPARANASE TO IDENTIFY AND
ITILE OF INVENTION: USE OF HEPARANASE COMPOUNDS
ORRESPONDENCES: 43
CORRESPONDENCE ADDRESS:
ADDRESSE: The Upjohn Company, Intellectual Property La
STREET: 301 Henrietta
CITY: Kalamazco
STATE: MI
COUNTRY: USA
ZIP: 49001
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PALENTIN Release #1.0, Version #1.25
SOFTWARE: PALENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/08207A
FILING DATE:
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Mismatches

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100.0%; Similarity 100.0%; 2; Conservative 0

92;

Length

Score 486; DB 1; Pred. No. 5.1e-48;

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1 MKLCVTVLSLLMLVAAFCSPALSAPMGSDPPTACCFSYTARKLPRNFVVDYYETSSLCSQ 60

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Sequence 28, Application PC/TUS9408207

GENERAL INFORMATION:

APPLICANT: Hoogewerf, Arlene J.

APPLICANT: Ledbetter, Steven R.

TITLE OF INVENTION: USE OF HEPARANASE COMPOUNDS

TITLE OF INVENTION: ISOLATE ANTI-HEPARANASE COMPOUNDS

CORRESPONDENCE ADDRESS:

ADDRESSEE: The Upjohn Company, Intellectual Property Law

STREET: 301 Henrietta

CITY: Kalamazoo
                                                                                                                                                                                                                                                                                           Length 92;
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Pred. No. 5.1e-48;
Mismatches 0;
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/08207
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
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      ATTORNEY/AGENT INFORMATION:
NAME: Jameson, William G.
REGISTRATION NUMBER: 27,199
REFERENCE/DOCKET NUMBER: 4731.1 CI
TELECOMMUNICATION INFORMATION:
TELEFHONE: 616/385-7561
TELEFAX: 616/385-6897
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REGISTRATION NUMBER: 27,199
REFERENCE/DOCKET NUMBER: 4731
TELECOMMUNICATION INFORMATION:
TELEPHONE: 616/385-7561
TELEFAX: 616/385-6897
TELEX: 224401
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
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CITY: Kalam
STATE: MI
COUNTRY: US
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                                                                                                                           APPLICANT: The General Hospital Corporation
TITLE OF INVENTION: MCP-4 AND MCP-5: NOVEL CHEMOKINES
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Clark & Elbing LLP
STREET: 176 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 486; DB 1;
Pred. No. 5.1e-48;
; Mismatches 0;
61 PAVVFQTKRSKQVCADPSESWVQEYVYDLELN 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 PAVVFQTKRSKQVCADPSESWVQEYVYDLELN 92
                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/TTCT / TLING DATE:
CLASSTO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 4
PCT-US98-04002-10
; Sequence 10, Application PC/TUS9804002
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/027,128
FILING DATE: 30-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAME: Bieker-Brady, Kristina
REGISTRATION NUMBER: 39,109
REFERENCE/DOCKET NUMBER: 00786/2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-428-0200
TELEFAX: 617-428-7045
                                                                                                  Sequence 10, Application PC/TUS9717900 GENERAL INFORMATION:
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 92 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: protein
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92; Conserva
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PCT-US97-17900-10
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FILING DATE:
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US-08-136-117-28
US-08-136-117-28

Sequence 28, Application US/08136117
GENERAL INFORMATION:
APPLICANT: Hoogewerf, Arlene J.
APPLICANT: Ledbetter, Steven R.
TITLE OF INVENTION: USE OF HEPARANASE TO IDENTIFY AND TITLE OF INVENTION: ISOLATE ANTI-HEPARANASE COMPOUNDS
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: The Upjohn Company, Intellectual Property L:
CITY: Kalamazoo
CITY: Kalamazoo
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                                        POLYPEPTIDES CONTAINING
                                                                                                                                                                                 ZIP: 02140
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US98/04002
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
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Pred. No. 5.1e-48;
Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Herrmann, Steve
APPLICANT: Swanberg, Stephen
TITLE OF INVENTION: CHEMOKINE DOMAINS
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Insititute, Inc.
STREET: 87 CambridgePark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 49001
ZOMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/136,117
                                                                                                                                                                                                                                                                                                                                                                                             P-41,323
ER: GI5291
                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION.

NAME: Sprunger, Suzanne
REGISTRATION NUMBER: P-41,323
REFERENCE/DOCKET NUMBER: G152
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8284
TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               th 100.0%; Similarity 100.0%; F 92; Conservative 0;
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CITY: Ca-
STATE: M
COUNTRY:
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Matches 92
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GENERAL INFORMATION:
APPLICANT: Coleman, Roger
APPLICANT: Bandman, Olga
APPLICANT: Wilde, Craig G.
TITLE OF INVENTION: NEW CHEMOKINES EXPRESSED IN PANCREAS
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                           Indels
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100.0%; Pred. No. 5.1e-48;
tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Incyte Pharmaceuticals, Inc
CLASSIFICATION: 514

ATTORNEY/AGENT INFORMATION:
NAME: Jameson, William G.
REGISTRATION NUMBER: 27,199
REFERENCE/DOCKET NUMBER: 4731.1 CP
TELECOMMUNICATION INFORMATION:
TELEFHONE: 616/385-7561
TELEX: 224401
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
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33,954
33,954
T. PF-0027 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/390,740B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0195
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRIES 94304

ZIP: 94304

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette

MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3174 Porter Drive
Palo Alto
                                                                                                                                      ID NO:
ACTERISTICS:
A: 92 amino acids
IYPE: amino acid
STRANDEDNESS: 8:
TOPOLOGY: 1:
OLECULE
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TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                          Conservative
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OPERATING SYSTEM:
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MOLECULE TYPE: po
IMMEDIATE SOURCE:
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US-08-136-117-28
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CLONE: M
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STATE:
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61 PAVVFQTKRSKQVCADPSESWVQEYVYDLELN 92
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US-08-927-939-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapiens
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92; Conserv
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                                                                   RESULT 8
US-08-927-939-20
                                                                                                                                                                                                                                                                                                                                             92;
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                                                                                                                                                                                                                                      SEQ ID NO 20
LENGTH: 92
   61
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Coleman, Roger
APPLICANT: Wilde, Craig G.
APPLICANT: Seilhamer, Jeffrey J.
TITLE OF INVENTION: A NEW CHEMOKINE EXPRESSED IN FETAL SPLEEN,
TITLE OF INVENTION: ITS PRODUCTION AND USES
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Pred. No. 5.1e-48;
Mismatches 0;
                        Score 486; DB 7;
Pred. No. 5.1e-48
Mismatches 0
                                                                                                                                92
                                                                                                                             COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,123
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/375,346
FILING DATE: 19-JAN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: LUTHER, BARBARA J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0026 US
TELEPHONE: (415) 855-0555
                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: INCYTE PHARMACEUTICALS, STREET: 3330 HILLVIEW AVENUE CITY: PALO ALTO STATE: CALIFORNIA
                                                                                                                                                                                                       US-08-467-123-4; Sequence 4, Application US/08467123; GENERAL INFORMATION:
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100.0%;
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855-0572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 92 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compati)
                                                   Conservative
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HYPOTHETICAL: NO
ANTI-SENSE: NO
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                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amino acid
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                                     Similarity 92; Conser
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ORIGINAL SOURCE:
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APPLICANT:
APPLICANT:
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US-08-390-740B-7
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Sequence 20, Application US/08927939A

GENERAL INFORMATION:

APPLICANT: Grainger, David J.

APPLICANT: Tatalick, Lauren Marie

TITLE OF INVENTION: Compounds and methods to inhibit or

TITLE OF INVENTION: augment an inflammatory response.

FILE REFERENCE: 1543.001US1

CURRENT APPLICATION NUMBER: US/08/927,939A

CURRENT FILING DATE: 1997-11-09

NUMBER OF SEQ ID NOS: 85

SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                              methods to inhibit or
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100.0%; Pred. No. 5.1e-48;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                              Score 486; DB 13;
Pred. No. 5.1e-48;
; Mismatches 0;
Sequence 20, Application US/08927939; GENERAL INFORMATION:
APPLICANT: Grainger, David J.
APPLICANT: Tatalick, Lauen Marie
TITLE OF INVENTION: Compounds and methods to in
TITLE OF INVENTION: augment an inflammatory re;
FILE REFERENCE: 295.022US1
CURRENT APPLICATION NUMBER: US/08/927,939
CURRENT FILING DATE: 1997-09-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: FastSEQ for Windows Version 3.0
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GENERAL INFORMATION:
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INFORMATION FOR SEQUENCE CHARA
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GENERAL INFORMATION:
APPLICANT: Hedrick, Joseph A.
APPLICANT: Wang, Luquan
APPLICANT: Zlotnik, Albert
APPLICANT: Zlotnik, Albert
APPLICANT: Greene, Jonathan R.
TITLE OF INVENTION: Mammalian Chemokines; Related Reagents
TITLE OF INVENTION: and Methods
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                         CHEMOKINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 486; DB 13;
Pred. No. 5.1e-48;
0; Mismatches 0;
                                                                                                                                                                      COUNTRY: USA
ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/940,687
FILING DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/027,128
FILING DATE: 30-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAME: Bieker-Brady, Kristina
REGISTRATION NUMBER: 39,109
REFERENCE/DOCKET NUMBER: 00786/293002
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-428-7045
APPLICANT: Luster, Andrew D.
APPLICANT: Garcia-Zepeda, Eduardo A.
APPLICANT: Sarafi, Mindy N.
TITLE OF INVENTION: MCP-4 AND MCP-5: NOVEL CH.
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Clark & Elbing LLP
STREET: 176 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9
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901 California Avenue
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SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
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STATE: California
COUNTRY: USA
ZIP: 94304-1104
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                                                                                                   ADDRESSEE: ClassTREET: 176 FOR CITY: Boston STATE: MA COUNTRY: USA ZIP: 02110
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SCULE TYPE:
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-09-023-092-8
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Best Local 9
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Pred. No. 5.1e-48;
; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,230
                                                                                                        APPLICATION NUMBER: US/09/023,092
FILING DATE: 13-FEB-1998
CLASSIFICATION: 536
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0,

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                        DX0780
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3R: 97-23
                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Ching, Edwin P.
REGISTRATION NUMBER: 34,090
REFERENCE/DOCKET NUMBER: DX078
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650)852-9196
TELEFAX: (650)496-1200
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3, Application US/09081230
GENERAL INFORMATION:
APPLICANT: Sheppard, Paul
TITLE OF INVENTION: HUMAN CHEMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Lingenfelter, Susan E
REGISTRATION NUMBER: 41,156
REFERENCE/DOCKET NUMBER: 97-2.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-442-6675
TELEFAX: 206-442-6678
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ADDRESSEE: ZymoGenetics
STREET: 1201 Eastlake Ave.
CITY: Seattle
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Similarity 100.0%;
92; Conservative 0
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
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CHARACTERISTICS:
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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GY: linear
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                                                                                                   Length 92;
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GENERAL INFORMATION:
APPLICANT: BANDMAN, OLGA
APPLICANT: COLEMAN, ROGER
APPLICANT: STUART, SUSAN G.
TITLE OF INVENTION: NEW CHEMOKINE EXPRESSED IN EOSINOPHILS
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92
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                                                                                                                                 Indels
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/113,705
                                                                                                Score 486; DB 14;
Pred. No. 5.1e-48;
Mismatches 0;
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larity 100.0%; Pred. No. 5.1e-48;
Conservative 0; Mismatches 0;
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/421,144
FILING DATE: 13-APR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33954
REFERENCE/DOCKET NUMBER: PF-0031
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
INFORMATION FOR SEQ ID NO: 4:
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                                                                                                 Similarity 100.0%; 92; Conservative
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LENGTH: 92 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                           single
                                                      protein
amino
          TYPE: amino acid
STRANDEDNESS: sin
TOPOLOGY: linear
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92; Conser
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                                                    ; MOLECULE TYPE:
US-09-081-230-3
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US-09-113-705-4
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US-09-113-705-4
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            Sequence 7, Application US/09113705A
GENERAL INFORMATION:
APPLICANT: Bandman, Olga
APPLICANT: Coleman, Roger
APPLICANT: Stuart, Susan G.
TITLE OF INVENTION: NEW CHEMOKINE EXPRESSED IN EOSINOPHILS
FILE REFERENCE: PF-0031-1 DIV
CURRENT APPLICATION NUMBER: US/09/113,705A
CURRENT FILING DATE: 1998-07-10
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 3.0
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; GENERAL INFORMATION:
; APPLICANT: Grainger, David J.
; APPLICANT: Tatalick, Lauen Marie
; APPLICANT: Tatalick, Lauen Marie
; APPLICANT: Tatalick, Lauen Marie
; APPLICANT: Kanaly, Suzanne T.
; TITLE OF INVENTION: Compounds and methods to inhibit or TITLE OF INVENTION: augment an inflammatory response.
; FILE REFERENCE: 295.027US1
; CURRENT APPLICATION NUMBER: US/09/150,813
; CURRENT FILING DATE: 1997-09-11
; EARLIER APPLICATION NUMBER: US 08/927939
; EARLIER FILING DATE: 1997-09-11
; NUMBER OF SEQ ID NOS: 105
; SOFTWARE: FastSEQ for Windows Version 3.0
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5.1e-48;
nes 0;
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Pred. No. 5.1e-48;
); Mismatches 0;
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Mismatches
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100.0%; Pred. No. E
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larity 100.0%;
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US-09-113-705-7
                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
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92; Conserv
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92; Conserv
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ID NO 7
-09-113-705-7
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LENGTH: 92
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Best Local
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Search completed: December 16, 2003, 15:46:43 Job time : 168.41 secs

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1666, Ap
1051, Ap
1051, Ap
1051, Ap
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/cgn2_6/ptodata/2/paa/PCT_NEW_COMB.pep:*
/cgn2_6/ptodata/2/paa/US06_NEW_COMB.pep:*
/cgn2_6/ptodata/2/paa/US07_NEW_COMB.pep:*
/cgn2_6/ptodata/2/paa/US09_NEW_COMB.pep:*
/cgn2_6/ptodata/2/paa/US10_NEW_COMB.pep:*
/cgn2_6/ptodata/2/paa/US10_NEW_COMB.pep:*
/cgn2_6/ptodata/2/paa/US10_NEW_COMB.pep:*
                                  ; Search time 13.57; (without alignments) 366.280 Million cell
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US-10-332-038A-9
US-10-398-457-5
US-10-398-457-31
PCT-US03-21703-42
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RESULT 1
US-60-487-610-1666
US-60-487-610-1666
Sequence 1666, Application US/60487610
GENERAL INFORMATION:
APPLICANT: CARGILL, Michele
APPLICANT: CARGILL, Michele
APPLICANT: HUANG, HORGJIN
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001469
CURRENT APPLICATION NUMBER: US/60/487,610
CURRENT FILING DATE: 2003-07-17
NUMBER OF SEQ ID NOS: 97101
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1666
LENGTH: 92
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US-60-487-610-1677

Sequence 1677, Application US/60487610

GENERAL INFORMATION:

APPLICANT: CARGILL, Michele

APPLICANT: HUANG, Hongjin

TITLE OF INVENTION: LIVER FIBROSIS IN HEPATITIS C VIRUS-INFECTED SUBJITILE OF INVENTION: METHODS OF DETECTION AND USES THEREOF

TITLE OF INVENTION: METHODS OF DETECTION AND USES THEREOF

TITLE OF INVENTION WIMBER: US/60/487,610

CURRENT FILING DATE: 2003-07-17

NUMBER OF SEQ ID NOS: 97101

SOFTWARE: FastSEQ for Windows Version 4.0

LENGTH: 92
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Pred. No. 3.3e-50;
Mismatches 0;
US-10-398-457-10

US-10-398-457-3

US-10-398-457-7

US-10-474-794-277

US-10-474-794-279

US-10-474-794-279

US-10-474-794-279

US-10-474-794-279

US-10-332-038A-15

US-60-487-610-2476

US-60-487-610-2330

PCT-US03-28745-4

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US-10-646-770-18
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Pred. No. 3.3e-50;
Mismatches 0;
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PRIOR APPLICATION NUMBER: US 60/217,683
PRIOR FILING DATE: 2000-07-12
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin version 3.2
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COUNTRY: U.S.A.
ZIP: 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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GENERAL INFORMATION:
APPLICANT: WILLIAMS, TIMOTHY,
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| Sequence 1057, Application US/60485450
| Sequence 1057, Application US/60485450
| GENERAL INFORMATION:
| APPLICANT: CARAIL, Michele
| APPLICANT: CHANG, Sheng-Yung
| TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
| TITLE OF INVENTION: VIRUS-INFECTED SUBJECTS, METHODS OF DETECTION AND
| TITLE OF INVENTION: THEREOF
| TITLE OF INVENTIO
                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: CHANG, Sheng-Yung
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: RESPONSE TO INTERFERON TREATMENT IN HEPATITIS
TITLE OF INVENTION: VIRUS-INFECTED SUBJECTS, METHODS OF DETECTION
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CLOO1470
CURRENT APPLICATION NUMBER: US/60/485,450
CURRENT FILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 47859
SOFTWARE: Fast SEQ for Windows Version 4.0
SEQ ID NO 1051
LENGTH: 92
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                                                                                     Score 486; DB 7;
Pred. No. 3.3e-50
Mismatches 0
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Sequence 1051, Application US/60485450
GENERAL INFORMATION:
APPLICANT: CARGILL, Michele
APPLICANT: CHANG, Sheng-Yung
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Similarity 100.0%;
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                     ; ORGANISM: Homo sapiens
US-60-487-610-1677
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ORGANISM:
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TYPE: PRT ORGANISM:
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 Gaps
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                                                                                                                                                                                                                                                                             APPLICANT: Gryphon Therapeutics, Inc.
APPLICANT: Offord, Robin
APPLICANT: Gaertner, Hubert
APPLICANT: Hartley, Oliver
TITLE OF INVENTION: Chemokine Receptor Modulators, Production and Use FILE REFERENCE: 03504.271
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Pred. No. 3e-37;
Mismatches 0; Indels
Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
INT APPLICATION DATA:
APPLICATION NUMBER: US/10/622,134
FILING DATE: 18-Jul-2003
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: NIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR
CITY: ARLINGTON
STATE: VIRGINIA
                                                                                                           JOSE, PETER J.
GRIFFITHS-JOHNSON, DAVID A.
HSUAN, JOHN J.
TITLE OF INVENTION: CHEMOTACTIC CYTOKINE
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Length 72;
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                                                                                                                                                                                                                                                                                                                                  Score 359; DB 7; Length 72
Pred. No. 2.4e-35;
0; Mismatches 0; Indels
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PCT-US03-21703-42
; Sequence 42, Application PC/TUS0321703
; GENERAL INFORMATION:
; APPLICANT: Rupp, Fabio
; APPLICANT: Wang, Jianrui
; APPLICANT: Zhou, Ping
; APPLICANT: Wehrman, Tom
                                 Sequence 1052, Application US/60485450
Sequence 1052, Application US/60485450
GENERAL INFORMATION:
APPLICANT: CARGILL, Michele
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                                                                                                                                                                                                                                                                                                                                   73.9%;
78.3%;
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ilarity 95.7%;
Conservative
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ORGANISM: Escherichia Coli
-10-398-457-5
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                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                           SEQ ID NO 1052
LENGTH: 72
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US-10-398-457-5
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LENGTH: 69
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Best Local (
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US-60-487-610-1667

Sequence 1667, Application US/60487610

GENERAL INFORMATION:

APPLICANT: CARGILL, Michele

APPLICANT: HUANG, Hongjin

TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH

TITLE OF INVENTION: METHODS OF DETECTION AND USES THEREOF

TITLE REFERENCE: CL001469

CURRENT APPLICATION NUMBER: US/60/487,610

CURRENT FILING DATE: 2003-07-17

NUMBER OF SEQ ID NOS: 97101

SOFTWARE: FASESEQ for Windows Version 4.0

LENGTH: 72
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Pred. No. 8.7e-37;
0; Mismatches 0;
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Pred. No. 2.4e-35;
); Mismatches 0;
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/291,038
FILING DATE: 14-Apr-1999
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
APPLICATION NUMBER: US/08/615,232A
FILING DATE: 13-AUG-1996
APPLICATION NUMBER: GB 9318984
FILING DATE: 14-SEP-1993
APPLICATION NUMBER: GB 9408602
FILING DATE: 29-APR-1994
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                        FILING L... AARY J.

ATTORNEY/AGENT INFORMAY J.

REGISTRATION NUMBER: 32,955

REFERENCE/DOCKET NUMBER: 550-3:

TELECOMMUNICATION INFORMATION:

TELEPHONE: (703) 816-4000

TELEPHONE: (703) 816-4100
                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 10
                                                                                                                                                                                                                                                                               NCE CHARACTERISTICS:
LENGTH: 68 amino acids
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Similarity 100.0%;
68; Conservative (
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TELEFAX: (703) 8:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTI
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                                                                                                                                                                                                                                                                                                              TYPE: amino
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-10-622-134-10
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Best Local S
Matches 68
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Best Local S
Matches 72
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Sequence 5, Application US/10398457
GENERAL INFORMATION:
APPLICANT: APPLIED RESEARCH SYSTEMS ARS HOLDING N.V.
TITLE OF INVENTION: CHEMOKINES MUTANTS IN THE TREATMENT OF MULTIPLE SCLEROSIS FILE REFERENCE: WO465
CURRENT APPLICATION NUMBER: US/10/398,457
CURRENT FILING DATE: 2003-11-13
NUMBER OF SEQ ID NOS: 37
SOFTWARE: Patentin version 3.0
APPLICANT: CHANG, Sheng-Yung
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: RESPONSE TO INTERFERON TREATMENT IN HEPATITIS C
TITLE OF INVENTION: VIRUS-INFECTED SUBJECTS, METHODS OF DETECTION AND USES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CL001470
CURRENT APPLICATION NUMBER: US/60/485,450
CURRENT APPLICATION NUMBER: US/60/485,450
NUMBER OF SEQ ID NOS: 47859
SOFTWARE: FastSEQ for Windows Version 4.0
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SSULT 12
3-60-485-450-1058
Sequence 1058, Application US/60485450
GENERAL INFORMATION:
APPLICANT: CHANG, Sheng-Yung
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: RESPONSE TO INTERFERON TREATMENT IN HEPATITIS C
TITLE OF INVENTION: VIRUS-INFECTED SUBJECTS, METHODS OF DETECTION AND USES
TITLE OF INVENTION: THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 MKLCVTVLSLLMLVAAFCSPALSAPMGSDPPTACCFSYTARKLPRNFVVDYYETSSLCSO 60
Sequence 8, Application US/10332038A;
GENERAL INFORMATION:
APPLICANT: Gryphon Therapeutics, Inc.
APPLICANT: Offord, Robin
APPLICANT: Offord, Robin
APPLICANT: Gaertner, Hubert
APPLICANT: Gaertner, Hubert
APPLICANT: Hartley, Oliver
TITLE OF INVENTION: Chemokine Receptor Modulators, Production and Use
TITLE OF INVENTION: Chemokine Receptor Modulators, Production and Use
CURRENT APPLICATION NUMBER: US/10/332,038A
CURRENT FILING DATE: 2003-05-06
PRIOR FILING DATE: 2000-07-12
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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; Pred. No. 8.6e-26;
13; Mismatches 9;
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                                                                               PAVVFQTKRSKQVCADPSESWVQEYVYDLELN 92
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                                                                                                                                                                                                                                                      GENERAL INTERPLICANT: CANCEL APPLICANT: CHANG, CAPPLICANT: CHANG, CAPPLICANT: CHANG, CAPPLICANT: CHANG, CAPPLICANT: CHANG, CAPPLICANT: CHANG, CAPPLICANTION: RESPONSE INTITLE OF INVENTION: VIRUS-INFECTED CAPPLICANTION: THEREOF FILE REFERENCE: CLOO1470 CURRENT APPLICATION NUMBER: US/60/485,450 CURRENT FILING DATE: 2003-07-09 NUMBER OF SEQ ID NOS: 47859 CURRENT FIRESE FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ed. No. 7.86
Mismatches
                                                                                                      60 PGVIFLTKRGRQVCADPSEEWVQKYVSDLELS
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7; Conservative 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   52;
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CENGTH: 70
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Best Local
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Best Local
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Matches
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US-60-487-610-1678

Sequence 1678, Application US/60487610

GENERAL INFORMATION:
APPLICANT: CARGILL, Michele
APPLICANT: HUANG, HONGJin
TITLE OF INVENTION: LIVER FIBROSIS IN HEPATITIS C VIRUS-INFECTED SUB,
TITLE OF INVENTION: METHODS OF DETECTION AND USES THEREOF
TITLE OF INVENTION UMBER: US/60/487,610
CURRENT APPLICATION NUMBER: US/60/487,610
CURRENT FILING DATE: 2003-07-17
NUMBER OF SEQ ID NOS: 97101
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1678
LENGTH: 92
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1: Methods and Materials Relating to Novel Polype
S-B2 PCT
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Pred. No. 7.8e-28;
19; Mismatches 20;
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CURRENT APPLICATION NUMBER: PCT/US03/21703
CURRENT FILING DATE: 2003-07-09
PRIOR APPLICATION NUMBER: PCT/US02/29636
PRIOR FILING DATE: 2002-09-18
PRIOR FILING DATE: 2002-07-12
PRIOR FILING DATE: 2002-07-13
PRIOR APPLICATION NUMBER: PCT/US02/22858
PRIOR FILING DATE: 2002-07-19
PRIOR FILING DATE: 2002-03-28
PRIOR FILING DATE: 2002-03-28
PRIOR FILING DATE: 2001-07-21
PRIOR FILING DATE: 2001-07-21
PRIOR FILING DATE: 2002-11-18
PRIOR FILING DATE: 2000-12-22
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Similarity 56.5%;
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 APPLICANT: Wang, Zhiwe
APPLICANT: Tang, Y. To
TITLE OF INVENTION: Met
FILE REFERENCE: HYS-B2
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; Sequence 31, Application US/10398457
; GENERAL INFORMATION:
; APPLICANT: APFLIED RESEARCH SYSTEMS ARS HOLDING N.V.
; TITLE OF INVENTION: CHEMOKINES MUTANTS IN THE TREATMENT OF MULTIPLE SCLEROSIS
; FILE REFERENCE: W0465
; FILE REFERENCE: W0465
; CURRENT FILING DATE: 2003-11-13
; NUMBER OF SEQ ID NOS: 37
; SOFTWARE: Patentin version 3.0
; SEQ ID NO 31
; LENGTH: 70
; TYPE: PRT
; ORGANISM: Escherichia coli
US-10-398-457-31
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PCT-US03-28745-3

Sequence 3, Application PC/TUS0328745

GENERAL INFORMATION:

APPLICANT: CHEMOKINE THERAPEUTICS CORP.

TITLE OF INVENTION: DESIGN OF CHEMOKINE ANALOGS FOR THE TREATMENT OF IT THE OF INVENTION NUMBER: PCT/US03/28745

CURRENT FILING DATE: 2003-09-11

NUMBER OF SEQ ID NOS: 1640

SOFTWARE: Patentin version 3.2

SEQ ID NO 3

LENGTH: 69

TYPE: PRT

ORGANISM: Artificial Sequence

FEATURE:

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Sequence 32, Appl Sequence 32, Appl Sequence 330, App Sequence 330, Appl Sequence 3, Appli Sequence 53, Appli Sequence 54, Appli Sequence 55, Appli Sequence 55, Appli
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Pred. No. 6.2e-48;
Mismatches 0;
US-09-834-794A-32

US-10-260-270-3

US-10-223-081-330

US-10-223-081-330

US-10-223-082-330

US-10-223-088-330

US-10-223-088-330

US-10-223-088-330

US-10-223-088-330

US-10-223-088-330

US-10-223-088-330

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US-10-223-089-330

US-10-223-089-330

US-10-223-089-330

US-10-223-089-330

US-10-301-822-179

US-09-334-954A-53

US-09-334-954A-53

US-09-334-954A-53

US-09-334-954A-53

US-09-908-600-3

US-09-920-137A-6

US-10-137-655-6

US-10-137-655-6

US-10-16-233A-53

US-10-16-275-221

US-10-16-275-221
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US-08-927-939-20
; Sequence 20, Application US/08927939
; Publication No. US20010006640A1
; GENERAL INFORMATION:
; APPLICANT: Grainger, David J.
; APPLICANT: Tatalick, Lauen Marie
; TITLE OF INVENTION: Compounds and methods to in
; TITLE OF INVENTION: augment an inflammatory res
; FILE REFERENCE: 295.022US1
; CURRENT APPLICATION NUMBER: US/08/927,939
; CURRENT FILING DATE: 1997-09-11
; NUMBER OF SEQ ID NOS: 83
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 20
: LENGTH: 92
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      Homo sapiens
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        Query Match
Best Local S
Matches 92
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PAVVFQTKRSKQVCADPSESWVQEYVYDLELN

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RESULT 2
US-09-834-795A-31
; Sequence 31, Application US/09834795A
; Patent No. US20020076710A1
; GENERAL INFORMATION:

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                                                                                                                                                                                                                                                                                                                 100.0%; Score 486; DB 9;
100.0%; Pred. No. 6.2e-48;
tive 0; Mismatches 0;
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Pred. No. 6.2e-48;
); Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 3
US-09-834-794A-31
Sequence 31, Application US/09834794A
Publication No. US20030026777A1
GENERAL INFORMATION:
APPLICANT: Lawrence, Papsidero
APPLICANT: Lyn, Dyster
APPLICANT: Lyn, Dyster
APPLICANT: Jana, Frustaci
TITLE OF INVENTION: Detection and Treatment of
FILE REFERENCE: 3380/11127-US4
CURRENT APPLICATION NUMBER: US/09/834,794A
CURRENT APPLICATION NUMBER: 09/146,580
PRIOR FILING DATE: 1998-09-03
PRIOR FILING DATE: 1998-01-20
PRIOR FILING DATE: 1998-01-20
PRIOR FILING DATE: 1998-07-09
NUMBER OF SEQ ID NOS: 35
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                 92
APPLICANT: Lawrence, Papsidero
APPLICANT: Lyn, Dyster
APPLICANT: Jana, Frustaci
TITLE OF INVENTION: Detection and Treatment o:
FILE REFERENCE: 3380/11127-US3
CURRENT APPLICATION NUMBER: US/09/834,795A
CURRENT FILING DATE: 2001-04-12
PRIOR APPLICATION NUMBER: 09/146,580
PRIOR FILING DATE: 1998-03-03
PRIOR FILING DATE: 1998-01-20
PRIOR FILING DATE: 1998-01-20
PRIOR FILING DATE: 1998-07-09
NUMBER OF SEQ ID NOS: 35
SOFTWARE: Patentin version 3.0
SEQ ID NO 31
LENGTH: 92
                                                     and Treatment
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                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens -09-834-795A-31
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Matches 92
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US-09-920-137A-7

US-09-920-137A-7

Sequence 7, Application US/09920137A

Publication No. US20030049725A1

GENERAL INFORMATION:

APPLICANT: Coleman, Roger

APPLICANT: Wilde, Craig G.

TITLE OF INVENTION: NEW CHEMOKINES EXPRESSED IN PANCREAS

NUMBER OF SEQUENCES: 11

CORRESPONDENCE ADDRESS:

ADDRESSE: Incyte Pharmaceuticals, Inc.

STREET: 3174 Porter Drive

CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WESULI
US-10-137-655-7

Sequence 7, Application US/10137655

Publication No. US20030138917A1

GENERAL INFORMATION:

APPLICANT: Coleman, Roger

APPLICANT: Wilde, Craig G.

TITLE OF INVENTION:

NUMBER OF SEQUENCES: 11

CORRESPONDENCE ADDRESS:

ADDRESSEE: Incyte Pharmaceuticals, Inc.

STREET: 3174 Porter Drive

CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 92;
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Pred. No. 6.2e-48;
; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/920,137A
                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-OTELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Diskette
COMPUTER: IBM COMPATIBLE
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MOLECULE TYPE: peptide
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single
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OPERATING SYSTEM:
SOFTWARE: FASTSEQ
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MIP-1b
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ZIP: 94304
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; CLONE: M
US-09-920-137A-7
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Pred. No. 6.2e-48;
; Mismatches 0;
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APPLICANT: Coleman, Roger
Wilde, Craig C.
Seilhamer, Jeffrey J.
TITLE OF INVENTION: CHEMOKINE EXPRESSED IN I
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ZIP: 94304
ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/158,366
FILING DATE: 29-May-2002
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/467,123B
FILING DATE: 06-JUN-1995
APPLICATION NUMBER: US 08/375,346
FILING DATE: 19-JAN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals,
STREET: 3174 Porter Drive
CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PAVVFQTKRSKQVCADPSESWVQEYVYDLELN
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/137,6!
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-002
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
TVPE: amino acid
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Publication No. US20020142398A1
GENERAL INFORMATION:
APPLICANT: Coleman, Roger
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ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide
IMMEDIATE SOURCE:
LIBRARY: GenBank
CLONE: MIP-1b
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US-10-137-655-7
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Best Local
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Wilde, Craig G.
TITLE OF INVENTION: NEW CHEMOKINES EXPRESSED IN PANCREAS
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
CITY: Palo Alto
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                                                                                                                                                                                                                                                                                                                               Length
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                                                                                                                                                                                                                                                                                                                    Score 486; DB 14;
Pred. No. 6.2e-48;
Mismatches 0;
ATTORNEY/AGENT INFORMATION:

NAME: Billings, Lucy J.

REGISTRATION NUMBER: 36,749

REFERENCE/DOCKET NUMBER: PF-0026-1 DIV

TELECOMMUNICATION INFORMATION:

TELEPHONE: 415-555-0555

TELEFAX: 415-845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/057,275
FILING DATE: 25-Jan-2002
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/390,740A
FILING DATE: February 17, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33,954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61 PAVVFOTKRSKOVCADPSESWVQEYVYDLELN 92
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
INFORMATION FOR SEQ ID NO: 7:
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TOPOLOGY: linear
;
MOLECULE TYPE: peptide
;
SEQUENCE DESCRIPTION: SEQ ID NO:
US-10-158-366-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 7, Application US/10057275
Publication No. US20020155545A1
GENERAL INFORMATION:
                                                                                                                            INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 92 amino acids
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TOPOLOGY: linear
MOLECULE TYPE: peptide
IMMEDIATE SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
                                                                                                                                                                                          TYPE: amino acid STRANDEDNESS: sin
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|SLCSQ 60
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US-10-293-705-11
Sequence 11, Application US/10293705
Publication No. US20030083468A1
GENERAL INFORMATION:
APPLICANT: Wilde, Craig G.
APPLICANT: Hawkins, Phillip R.
APPLICANT: Bandman, Olga
APPLICANT: Bendman, Olga
APPLICANT: Boolly Broomer, Jeffrey J.
TITLE REFERENCE: PF-0024-3 CON
CURRENT APPLICATION NUMBER: US/10/293,705
CURRENT FILING DATE: 2002-11-12
PRIOR FILING DATE: 1998-12-09
PRIOR FILING DATE: 1998-12-09
PRIOR FILING DATE: 1994-11-29
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PERL Program
SEQ ID NO 11
                                                                             92
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Rossi, Devora L.
Bacon, Kevin B.
Schall, Thomas J.
Zlotnik, Albert
INVENTION: MAMMALIAN CX3C CHEMOKINE GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 486; DB 15;
Pred. No. 6.2e-48;
0; Mismatches 0;
                                                                          486; DB 14;
No. 6.2e-48;
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                                                                             Score
Pred.
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APPLICANT: Hardiman, Gerard T.
APPLICANT: Rossi, Devora L.
APPLICANT: Bacon, Kevin B.
APPLICANT: Bacon, Kevin B.
APPLICANT: Schall, Thomas J.
APPLICANT: Schall, Thomas J.
APPLICANT: Zlotnik, Albert
TITLE OF INVENTION: MAMMALIAN CX3C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            / NAME/KEY: misc feature
/ OTHER INFORMATION: GenBank ID No.
US-10-293-705-11
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Similarity 100.0%;
92; Conservative C
LIBRARY: GenBank
CLONE: MIP-1D
SEQUENCE DESCRIPTION:
US-10-057-275-7
                                                                                                      Conservative
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                                                                                          Similarity
                                                                                                    92;
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-771-023-1
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Best Local S
Matches 92
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Best Local
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US-09-7
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                                                                                                                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 461; DB 11;
Pred. No. 4.4e-45;
1; Mismatches 3;
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Publication No. US20020150992A1
GENERAL INFORMATION:
APPLICANT: Wei et al.
TITLE OF INVENTION: Chemokine Alpha-5
FILE REFERENCE: PF401D1
CURRENT APPLICATION NUMBER: US/10/114,482
CURRENT FILING DATE: 2002-04-03
PRIOR APPLICATION NUMBER: 09/195,106
PRIOR FILING DATE: 1998-11-18
PRIOR FILING DATE: 1997-11-21
NUMBER OF SEQ ID NOS: 22
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 PAVVFQTKRSKQVCADPSESWVQEYVYDLELN
                                  Institute
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US/09/771,023
                                                                                                                                                                                                                                                                                                                                                               US 08/590,828
                                                                                                                                                                                                                                                                                                                                                                                                                                                   DX0569K2
                                                                                                                                                                                                                                                                                                            08/786,068
                              ADDRESSEE: DNAX Research Inst. STREET: 901 California Avenue CITY: Palo Alto STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-852-9196
TELEFAX: 415-496-1200
                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/786
FILING DATE: 1997-01-21
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/FILING DATE: 24-JAN-1996
ATTORNEY/AGENT INFORMATION:
NAME: Ching, Edwin P.
REGISTRATION NUMBER: 34,09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94.9%;
92.4%;
                                                                                                                                                                                                                                      APPLICATION NUMBER: US/
FILING DATE: 2001-01-25
CLASSIFICATION:
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                CORRESPONDENCE ADDRESS:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
                                                                                              COUNTRY: USA
ZIP: 94304-1104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           / TOPOLOGY: li
/ MOLECULE TYPE:
US-09-771-023-11
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US-10-114-482-3
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APPLICANT: Grainger, David J.
TITLE OF INVENTION: Compounds and methods to inhibit or
TITLE OF INVENTION: augment an inflammatory response;
FILE REFERENCE: 295.022US1;
CURRENT APPLICATION NUMBER: US/08/927,939
CURRENT FILING DATE: 1997-09-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: FastSEQ for Windows Version 3.0
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Pred. No. 5.8e-35;
0; Mismatches 0;
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  Mismatches
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CURRENT APPLICATION NUMBER: US/09/195,457
CURRENT FILING DATE: 1998-11-18
PRIOR APPLICATION NUMBER: 08/470,323
PRIOR FILING DATE: 1995-06-06
PRIOR FILING DATE: 1994-09-14
PRIOR FILING DATE: 1994-09-14
PRIOR FILING DATE: 1993-09-14
PRIOR FILING DATE: 1993-09-14
PRIOR FILING DATE: 1993-09-14
PRIOR FILING DATE: 1993-09-14
PRIOR FILING DATE: 1994-04-29
NUMBER OF SEQ ID NOS: 11
                                                                                                                                                                                                                                                                                                                   CHEMOTACTIC CYTOKINE
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                                                                                                                                                                                                                                             APPLICANT: WILLIAMS, TIMOTHY J.
APPLICANT: JOSE, PETER J.
APPLICANT: GRIFFITHS-JOHNSON, DAVID A.
APPLICANT: HSUAN, JOHN J.
TITLE OF INVENTION: CHEMOTACTIC CYTOKI
FILE REFERENCE: 550-33
                                                                                                                                                                                                     Sequence 10, Application US/09195457
Patent No. US20020081623A1
GENERAL INFORMATION:
APPLICANT: WILLIAMS, TIMOTHY J.
APPLICANT: JOSE, PETER J.
APPLICANT: GRIFFITHS-JOHNSON, DAVII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         76.3%; Sc.
100.0%; Pr
0;
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 Conservative
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                                                                                                             61 EYVYDLELN 69
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US-09-195-457-10
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69;
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ENGTH: 93
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Best Local
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                                                                                                                                                                                                     RESULT 11
US-09-792-793A-30
is Sequence 30, Application US/09792793A
is Patent No. US20020168370A1
is GENERAL INFORMATION:
is APPLICANT: McDonald, John R.
APPLICANT: McOggins, Philip
TITLE OF INVENTION: MTHODS AND COMPOSITIONS FOR TREATING SECONDARY TISS
if TITLE OF INVENTION: OTHER INFLAMMATORY CONDITIONS AND DISORDERS
if TITLE OF INVENTION UMBER: US/09/792,793A
CURRENT APPLICATION NUMBER: US/09/792,793A
CURRENT FILING DATE: 2001-02-22
is SOFTWARE: PatentIN Ver. 2.0
is SEQ ID NO 30
is LENGTH: 69
in LENGTH: 69
                                                                     RESULT 12
US-10-375-209A-30

Sequence 30, Application US/10375209A

Publication No. US20030215421A1

GENERAL INFORMATION:

APPLICANT: McDonald, John R.

APPLICANT: Coggins, Philip

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR TREATING SECONDARY TISS

TITLE OF INVENTION: METHODS AND COMPOSITIONS AND DISORDERS

FILE REFERENCE: 25020-601E

CURRENT APPLICATION NUMBER: US/10/375,209A

CURRENT FILING DATE: 2003-02-24

NUMBER OF SEQ ID NOS: 93

SOFTWARE: Patentin Ver. 2.0

LENGTH: 69

TYPE: PRT
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                              17;
11;
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Pred. No. 2.1e-35;
0; Mismatches 0;
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2.1e-35;
          81.1%; Score 394; DB 14; 77.2%; Pred. No. 1.9e-37; ive 10; Mismatches 11.
                                                                                                                                     9
                                                                                                                                   Score 375;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                th 77.2%; Sc. Similarity 100.0%; P 69; Conservative 0;
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                                         Conservative
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                         Local Similarity
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         Query Match
Best Local S
Matches 71
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Best Local S
Matches 69
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Pred. No. 5.5e-29;
; Mismatches 18;
                                                                                                                                                                       5215-PCT-US
                                                                                                                                                                                                                                    RESULT 15
US-09-151-450-2
Sequence 2, Application US/09151450
Sequence 2, Application US/09151450
Sequence 2, Application US/09151450
Sequence 2, Application US/09151450
SPELCANT:
APPLICANT:
COUNTRY:
COUNTRY:
COUNTRY:
COUNTRY:
COUNTRY:
COUNTRY:
CONPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER:
CONPUTER: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/151,450
FILING DATE:
CLASSIFICATION NUMBER: US/09/151,450
FILING DATE:
APPLICATION NUMBER: US/09/151,450
FILING DATE:
APPLICATION NUMBER: US/09/151,450
FILING DATE:
ATTORNEY/AGENT NUMBER: US/09/151,450
FILING DATE:
APPLICATION NUMBER: US/09/151,450
FILING DATE:
ATTORNEY/AGENT NUMBER:
ATTORNEY/AGENT NUMBER:
ATTORNEY/AGENT NUMBER
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AAM52447
ABU52388
AAR36770
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                                                                   The lymphokine/cytokine-like proteins are associated with the inflammatory response and/or have mitogenic activities. Antigens raised to the proteins may be useful in detection and purification, especially in bioassays of various tumour cells or genetic defects in the inflammatory response.

(Note: Revised entry submitted to correct the patent number format of US Government-owned NTIS applications to prevent clashes with ongoing granted patent numbers. For further information please visit the Derwe web site at www.derwent.com/dwpi/updates/ntis_us.html.)
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Pred. No. 4.4e-49;
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cloning and hybridisation using mRNA
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The protein was produced by cloning into the expression vector
PYMIP300. The human MIP-lalpha sequence was derived from the
lambda gt10 cDNA clone hMIP1-13a, and the GAPDH promoter sequence,
alpha factor transcription terminator derived from plasmid pGAI1,
the construction of which is described in patent application

EP0324-274. Recombinantly produced MIP proteins have diagnostic and
therapeutic utility for detecting and treating infections, cancer,
myelopoietic dysfunction and autoimmune diseases. Although not
directly cytotoxic for WEH1 tumor cells, MIP-1 treated macrophages
chibited enhanced antibody-independent macrophage cytotoxicity for
tumour targets. MIP-1 treatment stimulated proliferation of mature
tissue macrophages; this effect was synergistic with both CSF-1 and
GM-CSF. Purified preparations of the recombinantly derived
MIP-lalpha peptide alone induced TNF and IL-6 in macrophages, but
MIP-lalpha peptide alone induced TNF and IL-6 in macrophages.
TNF-inducton by MIP-lapha to a significant degree. Other
bloactivity defined for native MIP-1 and recombinant MIP-lalpha
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The proteins may also be used to determine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Purified heparanases, prepared under reducing conditions and activated with transglutaminase, are given in AAR70786-804. Most are prepared by reverse transcription of mRNA from activated huma leukocytes, then cloning of the cDNA into pVL1392 baculovirus vector, and expression in Sf9 cells in the presence of reduced glutathione and dithiothreitol.

(Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                           detecting
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is the inhibition of proliferation of less differentiated erythropoietin IL-3 dependent hematopoietic progenitor cells (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                       inflammatory protein 1-beta; MIP-1-beta;
heparin; heparan sulfate; arthritis; restenosis;
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                                                        ; DB 13;
4.4e-49;
hes 0;
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4.4e-49;
hes 0;
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                                                        Score 486; D
Pred. No. 4.4
; Mismatches
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larity 100.0%;
Conservative (
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93US-0136117
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cancer; wound healing.
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; AAQ85368.
                                                                                                                                                                                standard;
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                                                                Local Similarity
                                        92 AA;
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13-OCT-1993;
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                                                                                                                                                                                                                                                        Macrophage :
heparanase;
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29-AUG-1995
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                                        Sequence
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N-PSDB;
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Score 486; D Pred. No. 4.4 ; Mismatches

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from HUMACT2A. This sequence a number of the production of a construct from HUMACT2A. This sequence is used in the production of a construct comprising an isolated polynucleotide encoding a chimeric polypeptide which comprises at least 1 chemokine polypeptide covalently attached to at least 1 heterologous polypeptide. By including a heterologous protein in the construction, the chimeric polypeptides will have longer and increased biological activity and can direct the chemokine to a particular site. The chimeric polypeptides can also be designed to inhibit or desensitise chemokine receptors. They can be used to affect inhibiting angiogenesis, for regeneration of bone, cartilage, ligament or tendon, for recruiting inflammatory or autoimmune disorders. They can also be used as vaccine adjuvants or to enhance the activity of antigen presenting cells and for treating or preventing HIV infection.

Neutralising antibodies binding to the chimeric polypeptide may also be useful therapeutics for both conditions associated with the chemokine corresponding to the chemokine domain of the chimeric polypeptide and also in the treatment of some forms of cancer where abnormal expression of the chimeric colling and complete the conditions associated with the chemokine corresponding to the chemokine domain of the chimeric polypeptide and content where abnormal expression
    9
                              1 MXLCVTVLSLLMLVAAFCSPALSAPMGSDPPTACCPSYTARKLPRNFVVDYYETSSLCSQ 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chemokine, MIP-1beta, chimeric, human, heterologous protein, inhibitor, receptor, chemotaxis, migratory cell, angiogenesis, bone, regeneration, cartilage, ligament, tendon, bone marrow, transplant, inflammation, autoimmune disorder, vaccine adjuvant, antigen presenting cell, cancer, HIV, human immunodeficiency virus, therapy, prevention.
MKLCVTVLSLLMLVAAFCSPALSAPMGSDPPTACCFSYTARKLPRNFVVDYYETSSLCSO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human chemokine MIP-lbeta domain protein fragment.
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4.4e-49;
                                                                                                                                                                                          PAVVFQTKRSKQVCADPSESWVQEYVYDLELN 92
                                                                                                                                             PAVVFQTKRSKQVCADPSESWVQEYVYDLELN
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Pred. No. 4
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                                                                                                                                                                                                                                                                                                                                                                                                          AAW76225 standard; Protein; 92
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100.0%;
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Best Local Similarity
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gene therapy

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for detection, diagnosis, drug screening

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This sequence represents a human beta chemokine, Act-2 which is used to describe a method in which a novel beta chemokine, ZCHEMO-8 is isolated. Altered levels of ZCHEMO-8 may be indicative of pathological conditions, including infections, cancer, myelopoietic disorders, autoimmune including infections, cancer, myelopoietic disorders, autoimmune calculated infections, cancer, myelopoietic disorders, autoimmune calculated in including infections. The ZCHEMO-8 polypeptides can be used, or or reduce the damage in ischaemic and reperfusion injuries, in a wound healing regime to stimulate an infiltration of immune cells (e.g. monocytes, neutrophils, T lymphocytes or basophils) to a wound site to facilitate healing. ZCHEMO may be used to mobilise progenitor cells from the marrow into the peripheral blood for transplants. ZCHEMO-8 polypeptides could be used to further define the role of chemokines in mediating suppression of HIV replication in CD4+ T-cells and limiting progression of HIV infection to AIDS. Use may be made of ZCHEMO-8 antagonists may have a beneficial therapeutic effect in diseases where the inhibition of activation of certain macrophages, neutrophils, basophils, Blymphocytes and/or T cells may be effective. Such diseases include autoimmune diseases e.g. multiple sclerosis, insulin-dependent diabetes and systemic lupus erythematosus, rheumatoid arthritis, allergies, asthma or arteriosclerosis. Also benefit may be derived from using ZCHEMO-8 antagonists for chronic inflammatory and infective diseases. Antagonists may be used to dampen or inactivate and infective diseases. Antagonists may be used to dampen or inactivate and infective diseases. The products can also be used
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                                                                                                                                                                                                                                                                                                                                                                                              Chemokine; ZCHEMO-8; human; pathological condition; infection; cancer; autoimmune disorder; immunodeficiency; myelopoietic; wound healing; transplant; progenitor cell; HIV infection; AIDS; chemotherapy; radiation therapy; T cell; macrophage activation inhibitor; B lymphocyte; chronic inflammatory disease; infective disease; diagnosis; detection; drug screening; gene therapy; Act-2.
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mation, asthma or infections
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Mismatches
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                                                                                                             PAVVFOTKRSKOVCADPSESWVOEYVY
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                                                                                                                                                                                                                                                standard; Protein;
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Conservative
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                                                                                                                                                                                                                                                                                                                                                             protein
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                                                                                                                                                                                                                                                                                                                                                                     Macrophage recruitment; chemokine derivative; MCP-1; osteoporosis; monocyte chemoattractant protein-1; inflammation; atherosclerosis; HIV; AIDS; stroke; psoriasis; autoimmune disease; hypertension; endotoxaemia; basophil-mediated disease; myocardial infarction; acute ischaemia; rheumatoid arthritis; contraception.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New peptide 3, amide and heterocyclic compounds and saccharide conjugates used for inhibiting chemokine induced activity and for treating e.g. stroke, vascular diseases, autoimmune diseases and tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         autoimmune diseases, atherosclerosis, osteoporosis, HIV infection and AIDS, psoriasis, inflammatory diseases, hypertension, basophil-mediated diseases, endotoxaemia, myocardial infarction, acute ischaemia and rheumatoid arthritis, and can be used to prevent strokes and as contraceptives. The coding sequences for the chemokines can be used in gene therapy for the same diseases, as well as in the production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention concerns the identification of a number of chemokines which can be used to produce derivatives, agonists and antagonists which are then useful in disease treatment. The chemokines include sequences AABIS785-BIS794, AABIS803-BIS813 and AABIS831-BIS848 These chemokine derivatives can be used to treat diseases such as
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                              92;
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                            Score 486; DB 20;
Pred. No. 4.4e-49;
; Mismatches 0;
                                                                                                                                                              PAVVFQTKRSKQVCADPSESWVQEYVYDLELN
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99US-0271192.
99US-0452406.
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B; AAA74885.
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                                          Similarity
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17-MAR-1999;
01-DEC-1999;
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Sequence
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Best Local
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92 AA;

Sequence

Selenoprotein; HIV; Ebola virus; cancer; immune system disorder

HIV_Nefl fusion protein #13

Homo sapiens.

US6303295-B1

(first entry)

03-JUL-2002

AAM52446;

62

AAM52446 standard; Protein;

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                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                     e.g. human
System
                                   Selenoprotein; HIV; Ebola virus; cancer; immune system disorder,
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   92;
                   Indels
                                                                                                                                                                                                                                                                                                                                                                                 New selenoprotein for use in detecting certain viruses, e
immunodeficiency virus (HIV) or Ebola, cancer and immune
disorders -
  , DB 21;
4.4e-49;
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 core 486; Dired. No. 4.4
                                                                      73-76; 140pp; English.
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Pred.
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 100.0%;
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95US-003112P
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        Similarity
92; Conserv
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01-SEP-1995
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The present invention relates to selenoproteins encoded in the genome of a virus, where the coding sequence of the selenoprotein is genetically engineered for expression in a nucleic acid construct. The invention also discloses a method for identifying selenoprotein coding sequences, for detecting certain viruses (e.g. HIV or Ebola), cancer and immune system disorders. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chemokine, MIP-lbeta, chimeric, human, heterologous protein, inhibitor, receptor, chemotaxis, migratory cell, angiogenesis, bone, regeneration, cartilage, ligament, tendon, bone marrow, transplant, inflammation, autoimmune disorder, vaccine adjuvant, antigen presenting cell, cancer, HIV, human immunodeficiency virus, therapy, prevention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                          e.g. human
                                                                                                                                                                                                                                                                   New selenoprotein for use in detecting certain viruses, e.g. hur
immunodeficiency virus (HIV) or Ebola, cancer and immune system
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100.0%; Pred. No. 4.4e-49;
sive 0; Mismatches 0;
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                                                                                                                                                                                       Ramanathan CS;
                                                                                                                                                                                                                                                                                                                                                    Disclosure, Columns 73-76; 140pp, English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PAVVFQTKRSKQVCADPSESWVQEYVYDLELN
                                                                                                                                           (UYGE-) UNIV GEORGIA RES FOUND INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW76223 standard; Protein; 331 AA
                                           96US-0679493.
                                                                                   95US-001203P
95US-003112P
                                                                                                                                                                                     Nadimpalli RG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                               14-JUL-1995;
01-SEP-1995;
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Synthetic.
                                       12-JUL-1996;
16-OCT-2001.
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Similarity
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                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                 This sequence represents a human chemokine MIP-1beta domain, isolated from cDNA clone MPB-X. This sequence is used in the production of a construct comprising an isolated polynucleotide encoding a chimeric polypeptide which comprises at least 1 chemokine polypeptide covalently attached to at least 1 heterologous polypeptide. By including a charached to at least 1 heterologous polypeptide. By including a chemokine construction, the chimeric polypeptides will have longer and increased biological activity and can direct the chemokine to a particular site. The chimeric polypeptides can also be designed to inhibit or desensitise to migratory cells, e.g. for to affect the chemotactic recruitment of migratory cells, e.g. for stimulating or inhibiting angiogenesis, for regeneration of bone, cartilage, ligament or tendon, for recruiting transplanted bone marrow cells to bone marrow, or for treating or preventing inflammatory or autoimmune disorders. They can also be used as vaccine adjuvants or to enhance the activity of antigen presenting cells and for treating or centained with the chemokine corresponding to the chimeric polypeptide and also in the treatment of some forms of the chimeric polypeptide and also in the treatment of some forms of the chimeric polypeptide and also in the treatment of some forms of the chimeric polypeptide.
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                                                                                                                                                                                                                                                                    New chimeric polypeptide(s) - comprise chemokine polypeptide covalently linked to heterologous polypeptide, used for, e.g chemotactic recruitment of migratory cells
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100.0%; Pred. No. 2.1e-48;
tive 0; Mismatches 0;
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"Chemokine domain"
                                           peptide"
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           Location/Qualifiers
                                  signal "signal
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                                                                                                                                               98WO-US04002
                                                                                                                                                                   97US-0808720
                                                                                                                                                                                         GENETICS INST INC.
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                                            /note= "
25..331
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N-PSDB; AAV56825.
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92; Conserv
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Best Local
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Matches
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ABP6521
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pathene, and for resultable conditions, and for resultable conditions, and for resultable conditions and process and process to hypoxia conditions, reperfusion injury, for monitoring the therapeutic treatment of a disease or physiology, condition, such as cancer, ischaemic conditions, reperfusion injury, retinopathy, neonatal stress, preeclapmsia, atherosclerosis, inflammatory retinopathy, neonatal stress, preeclapmsia, erythropoiesis or hair loss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to methods for identifying genes and proteins that are implicated in a specific disease or physiological condition. The method comprises comparing the transcriptome/proteome of a specialised cell type implicated in a disease or condition with that of a second specialised cell type, under two experimental conditions, and identifying a gene that is differentially regulated in the two specialised cell types under experimental conditions. ABV77873-ABV78116 and ABP65061-ABP65257 were identified using the methods of the invention. The coding sequences and proteins are useful for treating a disease in a patient, for manufacture of a medicament for treating hypoxia-regulated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9
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                                                              Cytostatic; vasotropic; tranquiliser; antiatherosclerotic; gene therapy antiinflammatory; vulnerary; gynecological; ophthalmological; vaccine; hypoxia; tumourigenesis; angiogenesis; apoptosis; cancer; ischaemic condition; reperfusion injury; retinopathy; neonatal stress; preeclapmsia; atherosclerosis; inflammatory condition; wound healing; inflammation; erythropoiesis; hair loss; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identifying a gene involved in disease for treating hypoxia-regulated conditions, comprises comparing the transcriptome/proteome of two cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 MKLCVTVLSLLMLVAAFCSPALSAPMGSDPPTACCFSYTARKLPRNFVVDYYETSSLCSQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Harris RA;
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9.9e-49;
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Pred. No. 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Krige D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 35; Page 401; 538pp; English
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Hypoxia-regulated protein #93
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08-FEB-2001; 2001GB-0003156.
25-OCT-2001; 2001GB-0025666.
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98.9%;
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                                                                                                                                                                                                                                                                              The present invention relates to selenoproteins encoded in the genoa virus, where the coding sequence of the selenoprotein is genetica engineered for expression in a nucleic acid construct. The inventio discloses a method for identifying selenoprotein coding sequences, detecting certain viruses (e.g. HIV or Ebola), cancer and immune sy disorders. The present sequence was used to illustrate the inventio
                                                                                                                                                                                                                             e.g. human
system
                                                    Ebola virus; cancer; immune system disorder.
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                                                                                                                                                                                                                            detecting certain viruses, or Ebola, cancer and immune
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                                                                                                                                                                                                                                                                                                                                                               Score 482; DB 23;
Pred. No. 1.3e-48;
); Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                       92
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Ramanathan
                                                                                                                                                                        RES FOUND INC
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2
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                                   fusion protein #14
                                                                                                                            96US-0679493
                                                                                                                                            95US-001203P
                                                                                                                                                                                                                                                                                                                                                                99.2%;
larity 98.9%;
Conservative
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                   entry)
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                                                                                                                                                                                         Nadimpalli
                                                                                                                                                                       GEORGIA
                                                    HIV;
                                                                                                                                                                                                          WPI; 2002-024734/03.
                  (first
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                                                    Selenoprotein;
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                                                                                                                                                                      (UYGE-) UNIV
                                                                       sapiens
                                                                                        US6303295-B1
                                                                                                                           12-JUL-1996;
                                                                                                                                            14-JUL-1995;
01-SEP-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sapiens
                                                                                                                                                                                                                                                               Disclosure;
                 03-JUL-2002
                                                                                                          6-OCT-2001
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                                   HIV
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Burgess CE;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kekuda R, Tchernev VT, Liu X, Spytek KA, Patturajan M, Burg
Vernet CAM, Li L, Gorman L, Malyankar UM, Boldog FL, Guo X;
Shenoy SG, Padigaru M, Taupier RJ, Miller CE, Casman SJ, Pe
Gangolli EA, Gusev V, Smithson G, Zerhusen BD, Gerlach V;
Pochart PF, Fernandes ER, Shimkets RA, Rastelli L, Spaderna
Larochelle WJ, Zhong M, Khramtsov NV, Voss EZ, Herrmann JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 202; 413pp; English
                       2001US-274322P.
2001US-274322P.
2001US-274322P.
2001US-275579P.
2001US-275601P.
2001US-275601P.
2001US-276000P.
2001US-27600P.
2001US-277339P.
2001US-378152P.
2001US-31231P.
2001US-313182P.
2001US-313182P.
2001US-313182P.
2001US-313182P.
2001US-313182P.
2001US-313182P.
2001US-313182P.
2001US-313182P.
2001US-31318403P.
2001US-322360P.
2001US-322360P.
2001US-332486P.
2001US-332486P.
2001US-332486P.
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N-PSDB; ABX70479.
08-MAR-2001;

08-MAR-2001;

09-MAR-2001;

13-MAR-2001;

13-MAR-2001;

13-MAR-2001;

14-MAR-2001;

20-MAR-2001;

20-MAR-2001;

20-MAR-2001;

21-MAR-2001;

21-MAR-2001;

22-MAR-2001;

22-MAR-2001;

23-MAR-2001;

24-MAR-2001;

25-MAR-2001;

27-MAR-2001;

27-MAR-2001;

27-MAR-2001;

27-MAR-2001;

28-MAR-2001;

27-MAR-2001;

27-AUG-2001;

27-AUG-2001;
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09-NOV-2001;
09-NOV-2001;
07-MAR-2002;
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Gangolli EA,
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                                                                                                                                                                                                                                                                                                                                                                       Lucodstream of a mammal comprises admin. of MIP-1 and/or MIP-2 or their analogues to induce such a reduction. The MIP-1, MIP-2 or analogue may be operatively linked to a carrier. The MIPs can be used to inhibit megakaryocytopoiesis to effect in vivo reduction of platelet nos. They can be used to treat disorders with excessively high platelet counts such as thrombocytosis, stroke, pulmonary embol and myeloproliferative disorders.

(Updated on 25-MAR-2003 to correct DN first.
                                                                                                                                                                                            et;
                            MKLCVTVLSLLMLVAAFCSPALSAPMGSDPPTACCFSYTARKLPRNFVVDYYETS
                                     Macrophage inflammatory protein; megakaryocytopoiesis; MIP-1; MI
thrombocythemia; reactive thrombocytosis; stroke; emboli; platel
myeloproliferative disorder.
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Pred. No. 6.5e-48;
0; Mismatches 1;
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Score 478; DB 24;
Pred. No. 3.8e-48;
; Mismatches 1
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                                                            PAVVFQTKRSKQVCADPSESWVQEYVYDLELN
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                                                                                                                                                                                                                                                                                                                                                     or
                                                                                                                                                                                                                                                                                                                                            Suppression of megakaryocytopoiesis macrophage inflammatory protein-1 o
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98.4%;
illarity 97.8%;
Conservative
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Similarity 98.9%;
1; Conservative
                                                                                                                  standard; protein;
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90; Conser
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29-SEP-1993
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The invention relates to an isolated polynucleotide (I) comprising one of 245 sequences (ABQ93288-ABQ93532). Treating a condition comprising administering to a mammalian subject a composition comprising the protein (II) encoded by (I) (ABP62809-ABP63053) or an antibody (III) to (II).

(I), (II) and (III) are useful for diagnostic evaluation of disorders.

(I) is useful for gene therapy of diseases and (II) can be used for therapeutic treatment. Diseases that may be treated include wound healing and tissue repair, burns, central nervous system disorders (e.g. Alzheimer's, Parkinson's, Huntington's and amylotrophic lateral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the printed / from WIPO
                                                                                                                                                                                                                                                                                                                            antiparkinsonian; immunostimulant; cytostatic; immunosuppressive; antidiabetic; antiallergic; gene therapy; wound healing; tissue repair; burn; central nervous system disorder; Alzheimer's disease; Parkinson's disease; immune disorder; autoimmune disorder; multiple sclerosis; diabetes; allergy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polynucleotide and polypeptides, useful for treatment and diagnosis of Alzheimer's, Parkinson's, Huntington's, amylotrophic lateral sclerosis, immune deficiencies, cancer, autoimmune disorders, multiple sclerosis, diabetes and allergies -
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Note: The sequence data for this patent did not form part of the specification, but was obtained in electronic format directly fat ftp.wipo.int/pub/published_pct_sequences.
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                                                                                                                                                                                                                                                                                                            Human; vulnerary; dermatological; neuroprotective; nootropic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PAVVFQTKRSKQVCADPSESWVQEYVYDLELN 92
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u C, Drmanac RT,
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96.7%;
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                                                                                                                 standard; Protein;
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